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(54) Title: POLYMORPHISMS AND NEW GENES IN THE REGION OF THE HUMAN HEMOCHROMATOSIS GENE

(57) Abstract

Polymorphic sites in the region surrounding the HFE gene are provided. These polymorphisms are useful as surrogate markers in diagnostic assays for hemochromatosis. Additionally, a fine structure map of the 1 megabase region surrounding the HFE gene is provided, along with 235 kb of DNA sequence and 8 loci corresponding to candidate genes within the 1 megabase region, and in the purification of related proteins.

Polymorphisms and New Genes in the Region of the Human Hemochromatosis Gene

BACKGROUND OF THE INVENTION

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Hereditary hemochromatosis (HH) is an inherited disorder of iron metabolism wherein the body accumulates excess iron. In symptomatic individuals, this excess iron leads to deleterious effects by being deposited in a variety of organs leading to their failure, and resulting in cirrhosis, diabetes, sterility, and other serious illnesses. The gene which is defective in this disease was

10   disclosed in copending U.S.S.N. 08/652,265.

Fine structure mapping of the region to which the gene responsible for HH, HFE (denoted HH or HFE in some publications), was mapped makes possible the identification of candidate sequences comprising the HFE gene, along with structural elements for regulation and expression and neighboring genes.

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A variety of techniques is available for fine structure mapping, including direct cDNA selection, exon-trapping, and genomic sample sequencing. The direct selection approach (Lovett et al. Proc. Natl. Acad. Sci. U.S.A. 88:9628-9623 (1991)) involves the hybridization of cDNA fragments to genomic DNA. This technique is extremely sensitive and capable of isolating portions of rare transcripts. Exon-trapping (Church et al. Nature Genetics 6:98-105 (1994)) recovers spliced introns

20   from in vivo expressed genomic DNA clones and produces candidate exons without requiring any prior knowledge of the target's gene expression. High-throughput genomic DNA sequencing with comparison of the sequence data to databases of expressed sequences has also been used, such as in the positional cloning of the Werner syndrome gene (Yu et al. Science 277:258-262 (1996)) and in cloning by homology of the second Alzheimer's disease gene on chromosome 1 (Levy-Lahad et al.

25   Science 269:973-977 (1995)).

HH is typically inherited as a recessive trait; in the current state of knowledge, homozygotes carrying two defective copies of the gene are most frequently affected by the disease. In addition, heterozygotes for the HFE gene are more susceptible to sporadic porphyria cutanea tarda and potentially other disorders (Roberts et al., Lancet 349:321-323 (1997). It is estimated that

30   approximately 10-15% of Caucasians carry one copy of the HFE gene mutation and that there are about one million homozygotes in the United States. HH, thus, represents one of the most common genetic disease mutations in Caucasian individuals. Although ultimately HH produces debilitating symptoms, the majority of homozygotes and heterozygotes have not been diagnosed.

The need for such diagnostics is documented, for example, in Barton, J.C. et al.

35   Nature Medicine 2:394-395 (1996); Finch, C.A. West J Med 153:323-325 (1990); McCusick, V. Mendelian Inheritance in Man pp. 1882-1887, 11th ed., (Johns Hopkins University Press, Baltimore (1994)); Report of a Joint World Health Organization/Hemochromatosis Foundation/French Hemochromatosis Association Meeting on the Prevention and Control of Hemochromatosis (1993); Edwards, C.Q. et al. New Engl J Med 328:1616-1620 (1993); Bacon, B.R. New Engl J Med 326:126-

127 (1992); Balan, V. et al. Gastroenterology 107:453-459 (1994); Phatak, P.D. et al. Arch Int Med 154:769-776 (1994).

A single mutation in the HFE gene, designated 24d1 in copending U.S.S.N. 08/630,912, gave rise to the majority of disease-causing chromosomes present in the population today.

5   This is referred to herein as the "common" or "ancestral" or "common ancestral" mutation. These terms are used interchangeably. It appears that about 80% to 90% of all HH patients carry at least one copy of the common ancestral mutation which is closely linked to specific alleles of certain genetic markers close to this ancestral HFE gene defect. These markers are, as a first approximation, in the allelic form in which they were present at the time the ancestral HFE mutation occurred. See, for

10   example, Simon, M. et al. Am J Hum Genet 41:89-105 (1987); Jazwinska, E.C. et al. Am J Hum Genet 53:242-257 (1993); Jazwinska, E.C. et al. Am J Hum Genet 56:428-433 (1995); Worwood, M. et al. Brit J Hematol 86:863-866 (1994); Summers, K.M. et al. Am J Hum Genet 45:41-48 (1989).

Several polymorphic markers in the HFE region have been described and shown to have alleles that are associated with HH disease. These markers include the published microsatellite

15   markers D6S258, D6S306 (Gyapay, G. et al. Nature Genetics 7:246-339 (1994)), D6S265 (Worwood, M. et al. Brit J Hematol 86:833-846 (1994)), D6S105 (Jazwinska, E.C. et al. Am J Hum Genet 53:242-257 (1993); Jazwinska, E.C. et al. Am J Hum Genet 56:428-433 (1995)), D6S1001 (Stone, C. et al. Hum Molec Genet 3:2043-2046 (1994)), D6S1260 (Raha-Chowdhury et al. Hum Molec Genet 4:1869-1874 (1995)) as well as additional microsatellite and single-nucleotide-polymorphism markers

20   disclosed in co-pending PCT application WO 96/06583, the disclosure of which is hereby incorporated by reference in its entirety. Additionally, copending U.S.S.N. 08/630,912 disclosed additional markers 24d2 and 24d7.

The symptoms of HH are often similar to those of other conditions, and the severe effects of the disease often do not appear immediately. Accordingly, it would be desirable to provide a

25   method to identify persons who may be destined to become symptomatic in order to intervene in time to prevent excessive tissue damage associated with iron overload. One reason for the lack of early diagnosis is the inadequacy of presently available diagnostic methods to ascertain which individuals are at risk, especially while such individuals are presymptomatic.

Although blood iron parameters can be used as a screening tool, a confirmed

30   diagnosis often employs liver biopsy which is undesirably invasive, costly, and carries a risk of mortality. Thus, there is a clear need for the development of an inexpensive and noninvasive diagnostic test for detection of homozygotes and heterozygotes in order to facilitate diagnosis in symptomatic individuals, provide presymptomatic detection to guide intervention in order to prevent organ damage, and for identification of heterozygote carriers.

35   Furthermore, a need exists for both methods for fine structure mapping and a fine structure map of the region of the chromosome to which the HH locus maps. This and other needs are addressed by the present invention.

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SUMMARY OF THE INVENTION

One aspect of the invention is an oligonucleotide comprising at least 8 to about 100 consecutive bases from the sequence of Figure 9, or the complement of the sequence, wherein the at least 8 to about 100 consecutive bases includes at least one polymorphic site of Table 1.

Another aspect of the invention is an oligonucleotide pair selected from the sequence of Figure 9 or its complement for amplification of a polymorphic site of Table 1.

Another aspect of the invention is an isolated nucleic acid molecule comprising about 100 consecutive bases to about 235 kb substantially identical to the sequence of Figure 9, wherein the DNA molecule comprises at least one polymorphic site of Table 1.

Another aspect of the invention is a method to determine the presence or absence of the common hereditary hemochromatosis (HFE) gene mutation in an individual comprising:

providing DNA or RNA from the individual; and

assessing the DNA or RNA for the presence or absence of a haplotype of Table 1,

wherein, as a result, the absence of a haplotype of Table 1 indicates the likely absence of the HFE gene mutation in the genome of the individual and the presence of the haplotype indicates the likely presence of the HFE gene mutation in the genome of the individual.

Another aspect of the invention is a method to determine the presence or absence of the common hereditary hemochromatosis (HFE) gene mutation in an individual comprising:

providing DNA or RNA from the individual; and

assessing the DNA or RNA for the presence or absence of a genotype defined by a polymorphic allele of Table 1,

wherein, as a result, the absence of a genotype defined by a polymorphic allele of Table 1 indicates the likely absence of the HFE gene mutation in the genome of the individual and the presence of the genotype indicates the likely presence of the HFE gene mutation in the genome of the individual.

Another aspect of the invention is a culture of lymphoblastoid cells having the designation ATCC CRL-12371.

One aspect of the invention is an isolated nucleic acid sequence comprising a nucleic acid sequence substantially identical to BTF1.

A further aspect of the invention is an isolated nucleic acid sequence comprising a nucleic acid sequence substantially identical to BTF2.

A further aspect of the invention is an isolated nucleic acid sequence comprising a nucleic acid sequence substantially identical to BTF3.

A further aspect of the invention is an isolated nucleic acid sequence comprising a nucleic acid sequence substantially identical to BTF4.

A further aspect of the invention is an isolated nucleic acid sequence comprising a nucleic acid sequence substantially identical to BTF5.

A further aspect of the invention is an isolated nucleic acid sequence comprising a nucleic acid sequence substantially identical to NPT3.

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A further aspect of the invention is an isolated nucleic acid sequence comprising a nucleic acid sequence substantially identical to NPT4.

A further aspect of the invention is an isolated nucleic acid sequence comprising a nucleic acid sequence substantially identical to RoRet.

5   Additional aspects of the invention include nucleic acid sequences that are cDNAs, polypeptides encoded by the nucleic acids of the invention and antibodies specifically immunoreactive thereto, vectors comprising the nucleic acid sequences of the invention, and host cells stably transfected with the nucleic acids of the invention.

A further aspect of the invention is an isolated nucleic acid sequence comprising at 10   least 18 contiguous nucleotides substantially identical to at least 18 contiguous nucleotides of BTF1.

A further aspect of the invention is an isolated nucleic acid sequence comprising at least 18 contiguous nucleotides substantially identical to at least 18 contiguous nucleotides of BTF2.

A further aspect of the invention is an isolated nucleic acid sequence comprising at least 18 contiguous nucleotides substantially identical to at least 18 contiguous nucleotides of BTF3.

15   A further aspect of the invention is an isolated nucleic acid sequence comprising at least 18 contiguous nucleotides substantially identical to at least 18 contiguous nucleotides of BTF4.

A further aspect of the invention is an isolated nucleic acid sequence comprising at least 18 contiguous nucleotides substantially identical to at least 18 contiguous nucleotides of BTF5.

A further aspect of the invention is an isolated nucleic acid sequence comprising at 20   least 18 contiguous nucleotides substantially identical to at least 18 contiguous nucleotides of NPT3.

A further aspect of the invention is an isolated nucleic acid sequence comprising at least 18 contiguous nucleotides substantially identical to at least 18 contiguous nucleotides of NPT4.

A further aspect of the invention is an isolated nucleic acid sequence comprising at least 18 contiguous nucleotides substantially identical to at least 18 contiguous nucleotides of RoRet.

25   BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 depicts a combination genetic, physical and transcription map of the HFE gene region. The first line shows the relative positions of selected genetic markers that define the HFE region. The heavy bar below represents the YAC clone used in the direct selection experiment. The order and positions of the bacterial clones employed in the exon-trapping and sample sequencing is 30   indicated under the YAC. The thin bar under the bacterial clones represents the approximate locations of a subset of the expressed sequence fragments mapped to the contig. The thicker bars show the location of the cDNAs cloned. Two regions are bracketed; the butyrophilin family of genes (BTF), and the region where complete genomic sequencing was carried out.

Figure 2 is a schematic of the 250 kb of genomic sequence including the HFE gene. 35   Both the structure of the overall cDNA (top) and that corresponding to the coding regions (bottom), as well as the direction of transcription are shown. The positions of the histone genes, the zinc α-2 glycoprotein pseudogene, and the ESTs are also shown.

Figure 3 depicts an alignment of the predicted amino acid sequence of the BTF proteins. Sequences were aligned in a pair-wise fashion using CLUSTAL W (Thompson et al. Nucl. 40   Acids Res. 22:4673-4680) to deduce the most parsimonious arrangement. The asterisks under the

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alignment represent amino acids conserved in all 6 proteins; the "dots" represent conserved amino acids substitutions. Boxed are the regions within the proteins which correspond to three conserved motifs: 1) the B-G domain, 2) the transmembrane domain (TM), and 3) the B30-2 exon domain.

Figure 4, panel (A) depicts a Northern blot analysis of representative members of the two groups of BTF proteins, BTF1 and BTF5. BTF1 hybridized to all tissues on the blot as a major transcript at 2.9 kb and a minor one at 5.0 kb. BTF5 hybridized to several transcripts ranging between 4.0 and 3.1 kb and as a similar expression profile to BTF1. Autoradiography was for 24 hours. The β-actin hybridization demonstrated the variation in ploy (A)+ RNA between the lanes. Autoradiography was for 1 hour. In panel (B), RT-PCR analysis demonstrated that the expression of both genes was widespread. Included in the (+) lane are cDNA 21 and 44 as positive controls; the (-) lane represents the no-DNA control. Amplification using primers for the RFP gene (Isomura *et al. Nucleic Acid Res,* 20:5305-5310 (1992)) controlled for the integrity of the cDNA. All first strand cDNAs were checked for contaminating genomic DNA amplification by carrying out an identical experiment excluding the reverse transcriptase. In all cases, no amplification was obtained (data not shown).

Figure 5(A) depicts an alignment of the predicted amino acid sequence of the RoRet gene to the 52 kD Ro/SSA auto-antigen protein. The asterisks under the alignment represent conserved amino acids; the "dots" represent conserved amino acids substitutions. The putative DNA binding cysteine-rich domain and the B30-2 exon domain are boxed. Figure 5(B) depicts an alignment of the predicted amino acid sequence of the two novel putative sodium phosphate transport proteins to that of the NPT1.

Figure 6, panel (A) depicts a Northern blot analysis of the RoRet gene. The RoRet cDNA hybridized to 4 different transcripts, ranging from 7.1 kb to 2.2 kb. Autoradiography was performed for 4 days. The re-hybridization of the blot with a β-actin probe showed the variation in poly (A)+ RNA between the lanes. Autoradiography was for 1 hour. Panel (B) depicts RT-PCR analysis of the RoRet gene. Included in the (+) lane was a cDNA 27 positive control. Weak amplification of the correct size was observed in the small intestine, kidney and liver. The other tissues were negative as was the no DNA control lane (-). The RFP primers demonstrated the integrity of the cDNA. Panel (C) depicts Northern blot analysis of NPT3 and NPT4. NPT3 was expressed at high abundance in the heart and muscle as a single 7.2 kb transcript. Lesser amounts were found in the other tissues. The expression pattern of NPT4 was more restricted, being found only in the liver and kidney as a smear of transcripts ranging from 2.6 to 1.7 kb. Panel (D) depicts RT-PCR analysis of the NPT3 and NPT4 genes. Included in the (+) lane were the respective cDNA22E and 22B positive controls. The NPT3 gene was expressed as the proper size PCR fragment in kidney, liver, spleen and testis. A smaller fragment was detected in all tissues with the exception of the liver. The no DNA control lane (-) was negative. NPT4 was expressed as the proper size fragment in the small intestine, kidney, liver and testis. Larger and smaller size fragments were found in all other tissues with the exception of the brain. For both genes these different size fragments may indicate alternative splice events. The no DNA control lane (-) was negative. The RFP primers demonstrated the integrity of the cDNA.

Figure 7 depicts the sequences of cDNA 21 (BTF1), cDNA 29 (BTF3), cDNA 23 (BTF4), cDNA 44 (BTF5), cDNA 32 (BTF2), cDNA 27 (RoRet), cDNA 22B (NPT3), cDNA22E (NPT4).

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Figure 8 depicts the nucleotide sequence of approximately 235 kb in the HFE subregion from an unaffected individual.

Figure 9 depicts the nucleotide sequence of approximately 235 kb in the HFE subregion from an HH affected individual. Polymorphic sites in the HH affected individual determined by comparing a sequence of the corresponding region from an HH unaffected individual are listed and described in Table I.

DETAILED DESCRIPTION

A.   Definitions

Abbreviations for the twenty naturally occurring amino acids follow conventional usage. In the polypeptide notation used herein, the left-hand direction is the amino terminal direction and the right-hand direction is the carboxyl-terminal direction, in accordance with standard usage and convention. Similarly, unless specified otherwise, the left hand end of single-stranded polynucleotide sequences is the 5' end; the left hand direction of double-stranded polynucleotide sequences is referred to as the 5' direction. The direction of 5' to 3' addition of nascent RNA transcripts is referred to as the transcription direction; sequence regions on the DNA strand having the same sequence as the RNA and which are 5' to the 5' end of the RNA transcript are referred to as "upstream sequences"; sequence regions on the DNA strand having the same sequence as the RNA and which are 3' to the 3' end of the RNA transcript are referred to as "downstream sequences".

The term "nucleic acids", as used herein, refers to either DNA or RNA. "Nucleic acid sequence" or "polynucleotide sequence" refers to a single- or double-stranded polymer of deoxyribonucleotide or ribonucleotide bases read from the 5' to the 3' end. It includes both self-replicating plasmids, infectious polymers of DNA or RNA and nonfunctional DNA or RNA. The complement of any nucleic acid sequence of the invention is understood to be included in the definition of that sequence.

"Nucleic acid probes" may be DNA or RNA fragments. DNA fragments can be prepared, for example, by digesting plasmid DNA, or by use of PCR, or synthesized by either the phosphoramidite method described by Beaucage and Carruthers, _Tetrahedron Lett._ 22:1859-1862 (1981), or by the triester method according to Matteucci, _et al._, _J. Am. Chem. Soc._ 103:3185 (1981), both incorporated herein by reference. A double stranded fragment may then be obtained, if desired, by annealing the chemically synthesized single strands together under appropriate conditions or by synthesizing the complementary strand using DNA polymerase with an appropriate primer sequence. Where a specific sequence for a nucleic acid probe is given, it is understood that the complementary strand is also identified and included. The complementary strand will work equally well in situations where the target is a double-stranded nucleic acid.

The phrase "selectively hybridizing to" refers to a nucleic acid probe that hybridizes, duplexes or binds only to a particular target DNA or RNA sequence when the target sequences are present in a preparation of total cellular DNA or RNA. "Complementary" or "target" nucleic acid sequences refer to those nucleic acid sequences which selectively hybridize to a nucleic acid probe. Proper annealing conditions depend, for example, upon a probe's length, base composition, and the number of mismatches and their position on the probe, and must often be determined empirically. For

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discussions of nucleic acid probe design and annealing conditions, see, for example, Sambrook *et al.*, Molecular Cloning: a Laboratory Manual (2nd ed.), Vols. 1-3, Cold Spring Harbor Laboratory, (1989) or Current Protocols in Molecular Biology, F. Ausubel *et al.*, ed. Greene Publishing and Wiley-Interscience, New York (1987).

5   The phrase "nucleic acid sequence encoding" refers to a nucleic acid which directs the expression of a specific protein or peptide. The nucleic acid sequences include both the DNA strand sequence that is transcribed into RNA and the RNA sequence that is translated into protein. The nucleic acid sequences include both the full length nucleic acid sequences as well as non-full length sequences derived from the full length protein. It being further understood that the sequence

10   includes the degenerate codons of the native sequence or sequences which may be introduced to provide codon preference in a specific host cell.

The phrase "isolated" or "substantially pure" refers to nucleic acid preparations that lack at least one protein or nucleic acid normally associated with the nucleic acid in a host cell.

The phrase "expression cassette", refers to nucleotide sequences which are capable

15   of affecting expression of a structural gene in hosts compatible with such sequences. Such cassettes include at least promoters and optionally, transcription termination signals. Additional factors necessary or helpful in effecting expression may also be used as described herein.

The term "operably linked" as used herein refers to linkage of a promoter upstream from a DNA sequence such that the promoter mediates transcription of the DNA sequence.

20   The term "vector", refers to viral expression systems, autonomous self-replicating circular DNA (plasmids), and includes both expression and nonexpression plasmids. Where a recombinant microorganism or cell culture is described as hosting an "expression vector," this includes both extrachromosomal circular DNA and DNA that has been incorporated into the host chromosome(s). Where a vector is being maintained by a host cell, the vector may either be stably

25   replicated by the cells during mitosis as an autonomous structure, or is incorporated within the host's genome.

The term "gene" as used herein is intended to refer to a nucleic acid sequence which encodes a polypeptide. This definition includes various sequence polymorphisms, mutations, and/or sequence variants wherein such alterations do not affect the function of the gene product. The term

30   "gene" is intended to include not only coding sequences but also regulatory regions such as promoters, enhancers, and termination regions. The term further includes all introns and other DNA sequences spliced from the mRNA transcript, along with variants resulting from alternative splice sites.

The term "plasmid" refers to an autonomous circular DNA molecule capable of replication in a cell, and includes both the expression and nonexpression types. Where a recombinant

35   microorganism or cell culture is described as hosting an "expression plasmid", this includes both extrachromosomal circular DNA molecules and DNA that has been incorporated into the host chromosome(s). Where a plasmid is being maintained by a host cell, the plasmid is either being stably replicated by the cells during mitosis as an autonomous structure or is incorporated within the host's genome.

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The phrase "recombinant protein" or "recombinantly produced protein" refers to a peptide or protein produced using non-native cells that do not have an endogenous copy of DNA able to express the protein. The cells produce the protein because they have been genetically altered by the introduction of the appropriate nucleic acid sequence. The recombinant protein will not be found in association with proteins and other subcellular components normally associated with the cells producing the protein. The terms "protein" and "polypeptide" are used interchangeably herein.

The following terms are used to describe the sequence relationships between two or more nucleic acids or polynucleotides: "reference sequence", "comparison window", "sequence identity", "percentage of sequence identity", and "substantial identity". A "reference sequence" is a defined sequence used as a basis for a sequence comparison; a reference sequence may be a subset of a larger sequence, for example, as a segment of a full-length cDNA or gene sequence given in a sequence listing, or may comprise a complete cDNA or gene sequence.

Optimal alignment of sequences for aligning a comparison window may, for example, be conducted by the local homology algorithm of Smith and Waterman Adv. Appl. Math. 2:482 (1981), by the homology alignment algorithm of Needleman and Wunsch J. Mol. Biol. 48:443 (1970), by the search for similarity method of Pearson and Lipman Proc. Natl. Acad. Sci. U.S.A. 85:2444 (1988), or by computerized implementations of these algorithms (for example, GAP, BESTFIT, FASTA, and TFASTA in the Wisconsin Genetics Software Package Release 7.0, Genetics Computer Group, 575 Science Dr., Madison, WI).

The terms "substantial identity" or "substantial sequence identity" as applied to nucleic acid sequences and as used herein and denote a characteristic of a polynucleotide sequence, wherein the polynucleotide comprises a sequence that has at least 85 percent sequence identity, preferably at least 90 to 95 percent sequence identity, and more preferably at least 99 percent sequence identity as compared to a reference sequence over a comparison window of at least 20 nucleotide positions, frequently over a window of at least 25-50 nucleotides, wherein the percentage of sequence identity is calculated by comparing the reference sequence to the polynucleotide sequence which may include deletions or additions which total 20 percent or less of the reference sequence over the window of comparison. The reference sequence may be a subset of a larger sequence.

As applied to polypeptides, the terms "substantial identity" or "substantial sequence identity" mean that two peptide sequences, when optimally aligned, such as by the programs GAP or BESTFIT using default gap weights, share at least 80 percent sequence identity, preferably at least 90 percent sequence identity, more preferably at least 95 percent sequence identity or more. "Percentage amino acid identity" or "percentage amino acid sequence identity" refers to a comparison of the amino acids of two polypeptides which, when optimally aligned, have approximately the designated percentage of the same amino acids. For example, "95% amino acid identity" refers to a comparison of the amino acids of two polypeptides which when optimally aligned have 95% amino acid identity. Preferably, residue positions which are not identical differ by conservative amino acid substitutions. For example, the substitution of amino acids having similar chemical properties such as charge or polarity are not likely to effect the properties of a protein. Examples include glutamine for asparagine or glutamic acid for aspartic acid.

The phrase "substantially purified" or "isolated" when referring to a peptide or protein, means a chemical composition which is essentially free of other cellular components. It is preferably in a homogeneous state although it can be in either a dry or aqueous solution. Purity and homogeneity are typically determined using analytical chemistry techniques such as polyacrylamide gel electrophoresis or high performance liquid chromatography. A protein which is the predominant species present in a preparation is substantially purified. Generally, a substantially purified or isolated protein will comprise more than 80% of all macromolecular species present in the preparation. Preferably, the protein is purified to represent greater than 90% of all macromolecular species present. More preferably the protein is purified to greater than 95%, and most preferably the protein is purified to essential homogeneity, wherein other macromolecular species are not detected by conventional techniques.

The phrase "specifically binds to an antibody" or "specifically immunoreactive with", when referring to a protein or peptide, refers to a binding reaction which is determinative of the presence of the protein in the presence of a heterogeneous population of proteins and other biologies. Thus, under designated immunoassay conditions, the specified antibodies bind to a particular protein and do not bind in a significant amount to other proteins present in the sample. Specific binding to an antibody under such conditions may require an antibody that is selected for its specificity for a particular protein. A variety of immunoassay formats may be used to select antibodies specifically immunoreactive with a particular protein. For example, solid-phase ELISA immunoassays are routinely used to select monoclonal antibodies specifically immunoreactive with a protein. See Harlow and Lane (1988) Antibodies, a Laboratory Manual, Cold Spring Harbor Publications, New York, for a description of immunoassay formats and conditions that can be used to determine specific immunoreactivity.

As used herein, "EST" or "Expressed Sequence Tag " refers to a partial DNA or cDNA sequence of about 150 to 500, more preferably about 300, sequential nucleotides of a longer sequence obtained from a genomic or cDNA library prepared from a selected cell, cell type, tissue or tissue type, or organisms which longer sequence corresponds to an mRNA or a gene found in that library. An EST is generally DNA. One or more libraries made from a single tissue type typically provide at least 3000 different (i.e. unique) EST's and potentially the full complement of all possible EST's representing all possible cDNAs, e.g., 50,000 - 100,000 in an animal such as a human. (See, for example, Adams et al. Science 252:1651-1656 (1991)).

"Stringent" as used herein refers to hybridization and wash conditions of 50% formamide at 42°C. Other stringent hybridization conditions may also be selected. Generally, stringent conditions are selected to be about 5° C lower than the thermal melting point (Tm) for the specific sequence at a defined ionic strength and pH. The Tm is the temperature (under defined ionic strength and pH) at which 50% of the target sequence hybridizes to a perfectly matched probe. Typically, stringent conditions will be those in which the salt concentration is at least about 0.02 molar at pH 7 and the temperature is at least about 60°C. As other factors may significantly affect the stringency of hybridization, including, among others, base composition and size of the complementary strands, the presence of organic solvents and the extent of base mismatching, the combination of parameters is more important than the absolute measure of any one.

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B.   Transcript Map and New Genes near HH

The instant invention provides a fine structure map of the 1 megabase region surrounding the HFE gene. As part of that map the instant invention provides approximately 250 kb of DNA sequence of which about 235 kb are provided in Figure 8 and eight loci of particular interest
5   corresponding to candidate genes within the 1 megabase region. These loci are useful as genetic and physical markers for further mapping studies. Additionally, the eight cDNA sequences corresponding to those loci are useful, for example, for the isolation of other genes in putative gene families, the identification of homologs from other species, and as probes for diagnostic assays. In particular, isolated nucleic acid sequences of at least 18 nucleotides substantially identical to contiguous
10   nucleotides of a cDNA of the invention are useful as PCR primers. Typically, the PCR primer will be used as part of a pair of primers in a PCR reaction. Isolated nucleic acid sequences preferably comprising about 18-100 nucleotides, more preferably at least 18 nucleotides, substantially identical to contiguous nucleotides in a cDNA of the invention are useful in the design of PCR primers and probes for hybridization assays. Additionally, the proteins encoded by those cDNAs are useful in the
15   generation of antibodies for analysis of gene expression and in diagnostic assays, and in the purification of related proteins.

Thus, in one embodiment of the invention, a 235 kb sequence is provided for the HFE subregion within the 1 megabase region mapped. This sequence can serve as a reference in genetic or physical analysis of deletions, substitutions, and insertions in that region. Additionally, the sequence
20   information provides a resource for the further identification of new genes in that region. Thus, nucleic acid sequences substantially identically to the 235 kb sequence are also included in the scope of this invention.

In a further embodiment of the invention, a family of five genes, BTF1-5, is provided which are related by sequence homology to the milk protein butyrophilin (BT) (Figures 1, 3, and 7).
25   The predicted amino acid sequences of the proteins encoded by these genes are provided in Figure 3. These cDNAs are useful for the identification of further members of the BT family and to study regulation of expression of this family of genes. The proteins encoded by these cDNAs can be useful in the identification and isolation of ligands for the BT protein, and in the generation of agonists or antagonists of BT function. Nucleic acid sequences substantially identically to BTF1-5 and the proteins
30   encoded by them are also included in the scope of this invention, including allelic forms.

In a further embodiment of the invention, a novel gene RoRet is provided, which is related by sequence homology to the 52 kD Ro/SSA Lupus and Sjogren's syndrome autoantigen. This sequence is especially useful in the identification of other genes that may be involved in Lupus or Sjorgen's syndrome. The protein encoded by this cDNA can be useful in the identification and isolation
35   of ligands for the autoantigen, and in the generation of agonists or antagonists of the antigen. Nucleic acid sequences substantially identically to RoRet and the proteins encoded by them are also included in the scope of this invention.

In a further embodiment of the invention, two genes, NPT3 and NPT4, with structural homology to a type 1 sodium transport gene are provided. These cDNAs and the proteins expressed
40   by them are useful in determining the etiology of hypophosphatemia, along with being useful as probes

in the identification and isolation of further members of the gene family. Nucleic acid sequences substantially identically to th NPT1-like sequences and the proteins encoded by them are also included in the scope of this invention.

C.   Polymorphic Markers

The invention provides 397 new polymorphic sites in the region of the HFE gene. These polymorphisms are listed in Table 1. As described below, these polymorphisms were identified by comparison of the DNA sequence of an affected individual homozygous for the common ancestral HH mutation with that of an unaffected individual disclosed in copending U.S. 08/724,394.

Table 1. Polymorphic Sites in the HH Region

Base Location	Difference	Base Location	Difference
35-36	AC DEL	19755	G-A
841	T-C	19949	C-T
2662-2663	TT DEL	20085	C-T
3767	T-C	20366-20367	A INS
3829	C-G	20463	C-A
4925-4928	TAAA DEL	20841	A-T
5691	C-T	21059	A-T
5839	T-C	21117	A-G
6011	G-A	21837	A-C
6047	C-G	22293	A-C
6231	G-A	22786	C-A
6643	A DEL	23009	G-A
6698	T-C	24143	T-A
7186	T-C	26175	G-C
7273	G-A	26667	C-A
7545-7558	TCACACACCGATTGG DEL	26994	T-C
7672	G DEL	27838	G-T
7933	T-C	27861	T DEL
8746	T-G	28132	G-A
9115	G-A	29100	G-A
9823	G-A	29454-29457	TTTT DEL
10027	G-A	29787	T-G
10214	C-T	29825	A-C
10828	A-G	30009	T-C
10918	C-G	30177	A-G
10955	A-G	30400	A-G
11524	C-A	31059	T-A
11674	A-G	31280	C-T
11955	T-C	31749	C-T
12173-12175	TTT DEL	32040	C-G
13304	G-A	32556-32559	TGTG DEL
13455	G-A	33017	T-G
14416-14417	A INS	33026	T DEL
14998	C-T	34434	C-T
15564	T-C	35179	A-C
15887	A-G	35695	G-A
15904-15919	CCAAACTGATCTTTGA DEL	35702	G-A
16019	T DEL	35983	A-G
16211	A-T	37411	A-G
17461	A-G	38526	C-T

Base Location	Difference	Base Location	Difference
40431	C-A	72688	C-G
42054-42055	TT DEL	75323-75324	T INS
43783-43784	TTTT INS	75887	G-C
45120	C DEL	77519	T-C
45567	A-C	77749	G-A
46601	A-T	77908	T-C
47255	C-G	78385	C-G
47758	C-A	78592-78593	AG INS
47994	G-C	80189	T-G
48440	G-A	80279	T DEL
48650	T-G	80989-80990	A INS
48680	A-G	81193	T-C
50240	C-T	81273	A DEL
50553	G-A	82166	G-A
50586	G-T	83847	T DEL
51322	G-C	84161-84162	CA-GG
51747	A-G	84533	A-G
52474	C-G	84638	T-G
52733	C-A	85526	T-G
52875	G-A	85705	G-T
53631-53637	TTTTTTT DEL	86984	T-C
53707	G-A	87655	T-C
54819	A-G	87713	A-C
55913	T-C	87892	C-T
56225	A-C	88192	T DEL
56510	T-C	88528	A-G
56566	G-A	89645	A-T
56618	A-T	89728	A-G
57815	A-G	90088	T-C
58011	T DEL	91193-91194	2209bp INS
58247-58248	T INS	91373	T-C
58926	C-G	91433-91434	A INS
59406	C-G	91747	G-A
59422	G-C	93625	T DEL
60221-60222	A INS	95116-95117	T INS
60656-60657	CA DEL	96315	G-A
61162	G-A	97981	A-G
61465	G-A	98351	T DEL
61607	A DEL	99249	C-T
61653	T-C	100094-100095	T INS
61794-61795	T INS	100647-100648	TTC INS
62061	G-C	100951	C-T
62362	T-G	101610	C-G
62732	C-G	102589	C-T
63364	G-A	103076-103077	TATATATATATATA INS
63430-63431	GT INS	103747	T-C
63754	C-T	105638	A-C
63785	A-C	107024	C-T
63870-63871	A INS	107322	C-T
64788	A-G	107858	C-G
64962	G-A	109019	A DEL
65891	C-T	109579	T DEL
66675	G-C	110021	C-A
67186-67187	ATT INS	111251	C-A
67746-67747	TT INS	111425	G-A
68259	T-C	112644	T-A
68836	T-C	113001	G-C
68976	C-G	113130	C-T
72508	T-G	114026	G-A

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Base Location	Difference	Base Location	Difference
114250	A DEL	176222	T-C
115217	C-G	176524	A-T
117995	G-A	176684	G-A
118874	A-G	176815	T-C
119470	T-C	177049	T-C
119646	G-T	177065	G-T
120853	C-T	178285	T-C
121582	G-A	178551-178552	CTTTTTTTTTTTTTT INS
123576	A-C	179114-179115	A INS
125581	C-T	179260	C-G
125970	G-T	179281	C-G
126197	A-G	180023	G-C
126672	A DEL	180430	T-C
126672	G-C	180773	T-C
128220-128221	A INS	180824	T-C
132569	C-T	181097	C-T
133572	A-C	181183	A-T
134064	T-G	182351	C-T
136999	G-A	183197	G-A
137784	C-T	183623	A-T
138903	G-A	183653	G-T
139159-139160	A INS	183657	T-G
140359	G-A	183795-183796	A INS
140898	C-T	184060	G-A
141313	C DEL	184993	G-A
141343	T-C	185918	A-G
142148	T-C	186036	T-C
142178	C-A	186506-186507	TAAC INS
142433-142434	ATAGA INS	186561-186568	TATTTATT DEL
143783	C-T	186690	G DEL
144090	C-T	186751	T-A
144220-144221	A INS	187221	A-G
144725	A-C	187260	A-G
145732-145733	AAAAAAAAAAAAAA INS	187444-187447	CTCT DEL
147016-147017	CG DEL	187831-187832	C INS
147021	G-T	188638	G-A
147536	T-G	188642	C-T
148936	T-A	189246	T-C
149061	T-C	190340	A-C
154341	A-T	190354	A-G
154588	G-A	190762	A-G
155464	G-A	191260	G-T
158574	C-G	193018-193019	AGAT INS
160007	C-T	193147	T-G
164348	A-T	193196-193197	C INS
164499	C-G	193499	C-T
166677-166678	AAAG INS	193738	C-G
167389	G-A	193984-193985	ACACACAC INS
168506-168507	AGGATGGTCT INS	194064	C-G
168515	T-C	194504	A DEL
169413-169414	AA INS	194734	G-A
170300-170301	TTGTTGTTGTTG INS	194890	A-C
170491	G-A	195404	G-A
173428	T-C	195693	A-T
173642	G-A	196205	G-A
173948	T-G	197424	C-T
175330	T-C	197513	C-T
175836	T-C	197670	G-A
176200	G-C	198055	C-A

Base Location	Difference	Base Locati n	Difference
198401	C-T	215947	C-A
198692	A-G	216232	A-G
198780	T DEL	217478	G-A
199030	T-G	219052	T-C
199933	C-T	219082-219083	ATATATATATATATATATAT INS
200027	G-A	219314	C-A
200439	T-A	219327	G-A
200452	A-G	219560	C-T
200472-200483	AATAATAATAAT DEL	219660	C-T
200559	A-T	219889	G-A
200745	A-G	220198	G-T
200919	T-A	220384	G-A
201816	C-T	220451-220452	CAAAAA INS
201861-201862	42bp INS	221363	G-A
202662	T-C	221645	G-A
202880	T-C	222119	T-C
204341	C-T	222358	A-G
204768	A-T	222367	A-C
205284	T-G	222686	A-G
207400	C-A	222959	T-C
208634	T-C	223270-223271	TT DEL
208718	T DEL	223283	T-C
208862	A-C	224964	T-C
209419-209420	TT DEL	225232	A-C
209802	G-A	225366-225367	TTTT INS
209944	C-G	225416	G-C
210299	A-G	225486	T-C
211142	G-A	226088	A-G
212072	G-A	228421	A-G
212146	T-C	230047	G-A
212379	G-A	230109	G-C
212637-212639	TCT DEL	230376	C-G
212696	T-C	230394	A-G
213042	T-A	231226	A-G
214192	A-G	231447	G-A
214529-214530	TTTTTTTTTTTT INS	231835	A-G
214549	T-C	232400-232402	AAA DEL
214795	C-T	232402-232403	G INS
214908	T-G	232515	T-C
214977	A-G	232703	G-T
215769	C-T	232750	A-G

* D6S2238 occurs at base 1. 24d1 occurs at base 41316. D6S2239 occurs at base 84841. D6S2241 occurs at base 235032

Table 2. Polymorphic Allele Frequencies

Location	Frequency of ancestral variant in random chromosomes	Frequency of unaffected variant in random chromosomes
232703	53%	47%
231835	53%	47%
230394	85%	15%
230376	25%	75%
230109	53%	47%
225486	45%	55%
225416	75%	25%
220198	43%	57%
219660	58%	42%

Locati n	Frequency of ancestral variant in random chromosomes	Frequency f unaffected variant in random chromosomes
219560	53%	47%
214977	65%	35%
214908	50%	50%
214795	24%	76%
214549	53%	47%
214192	65%	35%
210299	53%	47%
208862	80%	20%
208634	48%	52%
207400	25%	75%
205284	50%	50%
204341	53%	47%
202880	58%	42%
202662	98%	2%
200027	25%	75%
199030	58%	42%
198692	55%	45%
198401	55%	45%
198055	55%	45%
195693	60%	40%
195404	25%	75%
194890	55%	45%
175330	53%	47%
173948	83%	17%
173642	55%	45%
173428	80%	20%
168515	80%	20%
160007	18%	82%
149061	58%	42%
148936	82%	18%
147536	100%	0%
147021	46%	54%
141343	55%	45%
140359	55%	45%
138903	55%	45%
132569	81%	19%
125581	18%	82%
121582	80%	20%
120853	18%	82%
118874	85%	15%
115217	50%	50%
113130	40%	60%
113001	48%	52%
107858	48%	52%
103747	50%	50%
96315	25%	75%
91194	80%	20%
90088	75%	25%
89728	50%	50%
89645	50%	50%
88528	63%	37%
87892	75%	25%
87713	60%	40%
87655	50%	50%
86984	79%	21%
85705	50%	50%
85526	50%	50%

Location	Frequency of ancestral variant in random chromosomes	Frequency of unaffected variant in random chrom s mes
84638	50%	50%
84533	50%	50%
82166	78%	22%
81193	58%	42%
80189	50%	50%
78385	80%	20%
77908	88%	12%
68976	50%	50%
68259	51%	49%
66675	80%	20%
62732	50%	50%
62362	40%	60%
61653	48%	52%
61465	5%	95%
61162	60%	40%
53707	100%	0%
52875	50%	50%
52733	74%	26%
52474	47%	53%
50586	50%	50%
50553	50%	50%
50240	50%	50%
48680	53%	47%
48650	63%	37%
48440	50%	50%
47255	50%	50%
46601	53%	47%
45567	49%	51%
41316	5%	95%
40431	20%	80%
38526	23%	77%
37411	70%	30%
35983	5%	95%

These polymorphisms provide surrogate markers for use in diagnostic assays to detect the likely presence of the mutations 24d1 and/or 24d2, in preferably 24d1, in homozygotes or heterozygotes. Thus, for example, DNA or RNA from an individual is assessed for the presence or absence of a genotype defined by a polymorphic allele of Table 1, wherein, as a result, the absence of a genotype defined by a polymorphic allele of Table 1 indicates the likely absence of the HFE gene mutation in the genome of the individual and the presence of the genotype indicates the likely presence of the HFE gene mutation in the genome of the individual.

These markers may be used singly, in combination with each other, or with other polymorphic markers (such as those disclosed in co-pending PCT application WO 96/06583) in diagnostic assays for the likely presence of the HFE gene mutation in an individual. For example, any of the markers defined by the polymorphic sites of Table 1 can be used in diagnostic assays in combination with 24d1 or 24d2, or at least one of polymorphisms HHP-1, HHP-19, or HHP-29, or microsatellite repeat all les 19D9:205; 18B4:235; 1A2:239; 1E4:271; 24E2:245; 2B8:206; 3321-1:98; 4073-1:182; 4440-1:180; 4440-2:139; 731-1:177; 5091-1:148; 3216-1:221; 4072-2:170; 950-1:142; 950-2:164; 950-3:165; 950-4:128; 950-6:151; 950-8:137; 63-1:151; 63-2:113; 63-3:169; 65-1:206; 65-

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2:159; 68-1:167; 241-5:108; 241-29:113; 373-8:151; and 373-29:113, D6S258:199, D6S265:122, D6S105:124; D6S306:238; D6S464:206; and D6S1001:180.

Table 2 lists the frequency of about 100 of the alleles defined by the polymorphic sites of the invention in the general population. As is evident from the Table, certain of these alleles are present rarely in the general population. These polymorphisms are thus preferred as surrogate markers in diagnostic assays for the presence of a mutant HFE allele ("gene mutation") such as 24d1 or 24d2. Preferably, the frequency of the polymorphic allele used in the diagnostic assay in the general population is less than about 50%, more preferably less than about 25%, and most preferably less than about 5%. Thus, of the genotypes defined by the alleles listed in Table 2, polymorphisms occurring at base 35983 and base 61465 of Figure 1 are preferred.

It will be understood by those of skill in the art that because they were identified in an ancestral HH homozygote, the haplotypes defined by the polymorphic sites of Table 1 are predictive of the likely presence of the HFE gene mutation 24d1. Thus, for example, the likelihood of any affected individual having at least two or more of any of the polymorphic alleles defined by Table 1 is greater than that for any unaffected individual. Similarly, the likelihood of any affected individual having at least three or more of any of the polymorphic alleles defined by Table 1 is greater than that for any unaffected individual.

Thus, for example, in a diagnostic assay for the likely presence of the HFE gene mutation in the genome of the individual, DNA or RNA from the individual is assessed for the presence or absence of a haplotype of Table 1, wherein, as a result, the absence of a haplotype of Table 1 indicates the likely absence of the HFE gene mutation in the genome of the individual and the presence of the haplotype indicates the likely presence of the HFE gene mutation in the genome of the individual.

The markers defined by the polymorphic sites of Table 1 are additionally useful as markers for genetic analysis of the inheritance of certain HFE alleles and other genes which occur within the chromosomal region corresponding to the sequence of Figure 9 which include, for example, those disclosed in copending U.S.S.N. 08/724,394.

As the entire nucleotide sequence of the region is provided in Figure 9, it will be evident to those of ordinary skill in the art which sequences to use as primers or probes for detecting each polymorphism of interest. Thus, in some embodiments of the invention, the nucleotide sequences of the invention include at least one oligonucleotide pair selected from the sequence of Figure 9 or its complement for amplification of a polymorphic site of Table 1. Furthermore, in some embodiments of the invention a preferred hybridization probe is an oligonucleotide comprising at least 8 to about 100 consecutive bases from the sequence of Figure 9, or the complement of the sequence, wherein the at least 8 to about 100 consecutive bases includes at least one polymorphic site of Table 1. In some embodiments the polymorphic site is at base 35983 or base 61465.

It will also be appreciated that the nucleic acid sequences of the invention include isolated nucleic acid molecules comprising about 100 consecutive bases to about 235 kb substantially identical to the sequence of Figure 9, wherein the DNA molecule comprises at least one polymorphic

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site of Table 1. Such isolated DNA sequences are useful as primers, probes, or as the component of a kit in diagnostic assays for detecting the likely presence of the HFE gene mutation in an individual.

D. Nucleic Acid Based Screening

Individuals carrying polymorphic alleles of the invention may be detected at either the

5   DNA, the RNA, or the protein level using a variety of techniques that are well known in the art. The genomic DNA used for the diagnosis may be obtained from body cells, such as those present in peripheral blood, urine, saliva, bucca, surgical specimen, and autopsy specimens. The DNA may be used directly or may be amplified enzymatically *in vitro* through use of PCR (Saiki et al. Science 239:487-491 (1988)) or other *in vitro* amplification methods such as the ligase chain reaction (LCR)

10   (Wu and Wallace Genomics 4:560-569 (1989)), strand displacement amplification (SDA) (Walker et al. Proc. Natl. Acad. Sci. U.S.A. 89:392-396 (1992)), self-sustained sequence replication (3SR) (Fahy et al. PCR Methods Appl. 1:25-33 (1992)), prior to mutation analysis. The methodology for preparing nucleic acids in a form that is suitable for mutation detection is well known in the art.

The detection of polymorphisms in specific DNA sequences, such as in the region of

15   the HFE gene, can be accomplished by a variety of methods including, but not limited to, restriction-fragment-length-polymorphism detection based on allele-specific restriction-endonuclease cleavage (Kan and Dozy Lancet ii:910-912 (1978)), hybridization with allele-specific oligonucleotide probes (Wallace et al. Nucl Acids Res 6:3543-3557 (1978)), including immobilized oligonucleotides (Saiki et al. Proc. Natl. Acad. Sci. U.S.A. 86:6230-6234 (1989)) or oligonucleotide arrays (Maskos and Southern

20   Nucl Acids Res 21:2269-2270 (1993)), allele-specific PCR (Newton et al. Nucl Acids Res 17:2503-2516 (1989)), mismatch-repair detection (MRD) (Faham and Cox Genome Res 5:474-482 (1995)), binding of MutS protein (Wagner et al. Nucl Acids Res 23:3944-3948 (1995), denaturing-gradient gel electrophoresis (DGGE) (Fisher and Lerman et al. *Proc. Natl. Acad. Sci. U.S.A.* 80:1579-1583 (1983)), single-strand-conformation-polymorphism detection (Orita et al. Genomics 5:874-879 (1983)), RNAase

25   cleavage at mismatched base-pairs (Myers et al. Science 230:1242 (1985)), chemical (Cotton et al. Proc. Natl. Acad. Sci. U.S.A. 85:4397-4401 (1988)) or enzymatic (Youil et al. Proc. Natl. Acad. Sci. U.S.A. 92:87-91 (1995)) cleavage of heteroduplex DNA, methods based on allele specific primer extension (Syvänen et al. Genomics 8:684-692 (1990)), genetic bit analysis (GBA) (Nikiforov et al. Nucl Acids Res 22:4167-4175 (1994)), the oligonucleotide-ligation assay (OLA) (Landegren et al. Science

30   241:1077 (1988)), the allele-specific ligation chain reaction (LCR) (Barrany Proc. Natl. Acad. Sci. U.S.A. 88:189-193 (1991)), gap-LCR (Abravaya et al. Nucl Acids Res 23:675-682 (1995)), radioactive and/or fluorescent DNA sequencing using standard procedures well known in the art, and peptide nucleic acid (PNA) assays (Orum et al., Nucl. Acids Res. 21:5332-5356 (1993); Thiede et al., Nucl. Acids Res. 24:983-984 (1996)).

35   In addition to the genotypes defined by the polymorphisms of the invention, as described in co-pending PCT application WO 96/35802 published November 14, 1996, genotypes characterized by the presence of the alleles 19D9:205; 18B4:235; 1A2:239; 1E4:271; 24E2:245; 2B8:206; 3321-1:98 (denoted 3321-1:197 therein); 4073-1:182; 4440-1:180; 4440-2:139; 731-1:177; 5091-1:148; 3216-1:221; 4072-2:170 (denoted 4072-2:148 therein); 950-1:142; 950-2:164; 950-3:165;

40   950-4:128; 950-6:151; 950-8:137; 63-1:151; 63-2:113; 63-3:169; 65-1:206; 65-2:159; 68-1:167; 241-

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5:108; 241-29:113; 373-8:151; and 373-29:113, alleles D6S258:199, D6S265:122, D6S105:124, D6S306:238, D6S464:206; and D6S1001:180, and/or alleles associates with the HHP-1, the HHP-19 or HHP-29 single base-pair polymorphisms can also be used to assist in the identification of an individual whose genome contains 24d1 and/or 24d2. For example, the assessing step can be
5 performed by a process which comprises subjecting the DNA or RNA to amplification using oligonucleotide primers flanking a polymorphism of Table 1, and oligonucleotides flanking 24d1 and/or 24d2, oligonucleotide primers flanking at least one of the base-pair polymorphisms HHP-1, HHP-19, and HHP-29, oligonucleotide primers flanking at least one of the microsatellite repeat alleles, or oligonucleotide primers for any combination of polymorphisms or microsatellite repeat alleles thereof.
10   Oligonucleotides useful in diagnostic assays are typically at least 8 consecutive nucleotides in length, and may range upwards of 18 nucleotides in length to greater than 100 or more consecutive nucleotides. Such oligonucleotides can be derived from either the genomic DNA of Figure 8 or 9, or cDNA sequences derived therefrom, or may be synthesized.

             Additionally, the proteins encoded by such cDNAs are useful in the generation of
15 antibodies for analysis of gene expression and in diagnostic assays, and in the purification of related proteins.

E.   **General Methods**

             The nucleic acid compositions of this invention, whether RNA, cDNA, genomic DNA, or a hybrid of the various combinations, may be isolated from natural sources, including cloned DNA,
20 or may be synthesized *in vitro*. The nucleic acids claimed may be present in transformed or transfected whole cells, in a transformed or transfected cell lysate, or in a partially purified or substantially pure form.

             Techniques for nucleic acid manipulation of the nucleic acid sequences of the invention such as subcloning nucleic acid sequences encoding
25 polypeptides into expression vectors, labeling probes, DNA hybridization, and the like are described generally in Sambrook *et al.*, Molecular Cloning - a Laboratory Manual (2nd Ed.), Vol. 1-3, Cold Spring Harbor Laboratory, Cold Spring Harbor, New York, (1989), which is incorporated herein by reference. This manual is hereinafter referred to as "Sambrook *et al.*"

             There are various methods of isolating the nucleic acid sequences of the invention.
30 For example, DNA is isolated from a genomic or cDNA library using labeled oligonucleotide probes having sequences complementary to the sequences disclosed herein. Such probes can be used directly in hybridization assays. Alternatively probes can be designed for use in amplification techniques such as PCR.

             To prepare a cDNA library, mRNA is isolated from tissue such as heart or pancreas,
35 preferably a tissue wherein expression of the gene or gene family is likely to occur. cDNA is prepared from the mRNA and ligated into a recombinant vector. The vector is transfected into a recombinant host for propagation, screening and cloning. Methods for making and screening cDNA libraries are well known. See Gubler, U. and Hoffman, B.J. Gene 25:263-269 (1983) and Sambrook *et al.*

             For a genomic library, for example, the DNA is extracted from tissue and either
40 mechanically sheared or enzymatically digested to yield fragments of about 12-20 kb. Th fragments

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are then separated by gradient centrifugation from undesired sizes and are constructed in
bacteriophage lambda vectors. These vectors and phage are packaged *in vitro*, as described in
Sambrook, *et al.* Recombinant phage are analyzed by plaque hybridization as described in Benton
and Davis, Science 196:180-182 (1977). Colony hybridization is carried out as generally described in
M. Grunstein *et al.* Proc. Natl. Acad. Sci. USA. 72:3961-3965 (1975).

DNA of interest is identified in either cDNA or genomic libraries by its ability to hybridize
with nucleic acid probes, for example on Southern blots, and these DNA regions are isolated by
standard methods familiar to those of skill in the art. *See* Sambrook, *et al.*

In PCR techniques, oligonucleotide primers complementary to the two 3' borders of
the DNA region to be amplified are synthesized. The polymerase chain reaction is then carried out
using the two primers. See PCR Protocols: a Guide to Methods and Applications (Innis, M, Gelfand,
D., Sninsky, J. and White, T., eds.), Academic Press, San Diego (1990). Primers can be selected to
amplify the entire regions encoding a full-length sequence of interest or to amplify smaller DNA
segments as desired.

PCR can be used in a variety of protocols to isolate cDNA's encoding a sequence of
interest. In these protocols, appropriate primers and probes for amplifying DNA encoding a sequence
of interest are generated from analysis of the DNA sequences listed herein. Once such regions are
PCR-amplified, they can be sequenced and oligonucleotide probes can be prepared from sequence
obtained.

Oligonucleotides for use as primers or probes are chemically synthesized according to
the solid phase phosphoramidite triester method first described by Beaucage, S.L. and Carruthers,
M.H., Tetrahedron Lett., 22(20):1859-1862 (1981) using an automated synthesizer, as described in
Needham-VanDevanter, D.R., *et al.*, Nucleic Acids Res. 12:6159-6168 (1984). Purification of
oligonucleotides is by either native acrylamide gel electrophoresis or by anion-exchange HPLC as
described in Pearson, J.D. and Regnier, F.E., J. Chrom., 255:137-149 (1983). The sequence of the
synthetic oligonucleotide can be verified using the chemical degradation method of Maxam, A.M. and
Gilbert, W., in Grossman, L. and Moldave, D., eds. Academic Press, New York, Methods in
Enzymology 65:499-560 (1980).

1.   **Expression**

Once DNA encoding a sequence of interest is isolated and cloned, one can express
the encoded proteins in a variety of recombinantly engineered cells. It is expected that those of skill in
the art are knowledgeable in the numerous expression systems available for expression of DNA
encoding a sequence of interest. No attempt to describe in detail the various methods known for the
expression of proteins in prokaryotes or eukaryotes is made here.

In brief summary, the expression of natural or synthetic nucleic acids encoding a
sequence of interest will typically be achieved by operably linking the DNA or cDNA to a promoter
(which is either constitutive or inducible), followed by incorporation into an expression vector. The
vectors can be suitable for replication and integration in either prokaryotes or eukaryotes. Typical
expression vectors contain transcription and translation terminators, initiation sequences, and
promoters useful for regulation of the expression of polynucleotide sequence of interest. To obtain

high level expression of a cloned gene, it is desirable to construct expression plasmids which contain, at the minimum, a strong promoter to direct transcription, a ribosome binding site for translational initiation, and a transcription/translation terminator. The expression vectors may also comprise generic expression cassettes containing at least one independent terminator sequence, sequences permitting

5   replication of the plasmid in both eukaryotes and prokaryotes, i.e., shuttle vectors, and selection markers for both prokaryotic and eukaryotic systems. See Sambrook et al. Examples of expression of ATP-sensitive potassium channel proteins in both prokaryotic and eukaryotic systems are described below.

a.   Expression in Prokaryotes

10   A variety of procaryotic expression systems may be used to express the proteins of the invention. Examples include E. coli, Bacillus, Streptomyces, and the like.

It is preferred to construct expression plasmids which contain, at the minimum, a strong promoter to direct transcription, a ribosome binding site for translational initiation, and a transcription/translation terminator. Examples of regulatory regions suitable for this purpose in E. coli

15   are the promoter and operator region of the E. coli tryptophan biosynthetic pathway as described by Yanofsky, C., J. Bacteriol. 158:1018-1024 (1984) and the leftward promoter of phage lambda (Pλ) as described by Herskowitz, I. and Hagen, D., Ann. Rev. Genet. 14:399-445 (1980). The inclusion of selection markers in DNA vectors transformed in E. coli is also useful. Examples of such markers include genes specifying resistance to ampicillin, tetracycline, or chloramphenicol. See Sambrook et

20   al. for details concerning selection markers for use in E. coli.

To enhance proper folding of the expressed recombinant protein, during purification from E. coli, the expressed protein may first be denatured and then renatured. This can be accomplished by solubilizing the bacterially produced proteins in a chaotropic agent such as guanidine HCl and reducing all the cysteine residues with a reducing agent such as beta-mercaptoethanol. The

25   protein is then renatured, either by slow dialysis or by gel filtration. See U.S. Patent No. 4,511,503.

Detection of the expressed antigen is achieved by methods known in the art as radioimmunoassay, or Western blotting techniques or immunoprecipitation. Purification from E. coli can be achieved following procedures such as those described in U.S. Patent No. 4,511,503.

b.   Expression in Eukaryotes

30   A variety of eukaryotic expression systems such as yeast, insect cell lines, bird, fish, and mammalian cells, are known to those of skill in the art. As explained briefly below, a sequence of interest may be expressed in these eukaryotic systems.

Synthesis of heterologous proteins in yeast is well known. Methods in Yeast Genetics, Sherman, F., et al., Cold Spring Harbor Laboratory, (1982) is a well recognized work describing the

35   various methods available to produce the protein in yeast.

Suitable vectors usually hav   expression control sequences, such as promoters, including 3-phosphoglycerate kinase or other glycolytic enzymes, and an origin of replication, termination sequences and th like as d sired. For instance, suitable vectors are describ d in the literature (Botstein, et al., Gene 8:17-24 (1979); Broach, et al., Gene 8:121-133 (1979)).

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Two procedures are used in transforming yeast cells. In one case, yeast cells are first converted into protoplasts using zymolyase, lyticase or glusulase, followed by addition of DNA and polyethylene glycol (PEG). The PEG-treated protoplasts are then regenerated in a 3% agar medium under selective conditions. Details of this procedure are given in the papers by J.D. Beggs, <u>Nature</u>

5   (London) 275:104-109 (1978); and Hinnen, a., *et al.*, <u>Proc. Natl. Acad. Sci. U.S.A.</u> 75:1929-1933 (1978). The second procedure does not involve removal of the cell wall. Instead the cells are treated with lithium chloride or acetate and PEG and put on selective plates (Ito, H., *et al.*, <u>J. Bact.</u> 153:163-168 (1983)).

The proteins of the invention, once expressed, can be isolated from yeast by lysing the

10   cells and applying standard protein isolation techniques to the lysates. The monitoring of the purification process can be accomplished by using Western blot techniques or radioimmunoassay or other standard immunoassay techniques.

The sequences encoding the proteins of the invention can also be ligated to various expression vectors for use in transforming cell cultures of, for instance, mammalian, insect, bird or fish

15   origin. Illustrative of cell cultures useful for the production of the polypeptides are mammalian cells. Mammalian cell systems often will be in the form of monolayers of cells although mammalian cell suspensions may also be used. A number of suitable host cell lines capable of expressing intact proteins have been developed in the art, and include the HEK293, BHK21, and CHO cell lines, and various human cells such as COS cell lines, HeLa cells, myeloma cell lines, Jurkat cells, etc.

20   Expression vectors for these cells can include expression control sequences, such as an origin of replication, a promoter (*e.g.*, the CMV promoter, a HSV *tk* promoter or *pgk* (phosphoglycerate kinase) promoter), an enhancer (Queen *et al.* <u>Immunol. Rev.</u> 89:49 (1986)), and necessary processing information sites, such as ribosome binding sites, RNA splice sites, polyadenylation sites (*e.g.*, an SV40 large T Ag poly A addition site), and transcriptional terminator sequences. Other animal cells useful for

25   production of ATP-sensitive potassium channel proteins are available, for instance, from the American Type Culture Collection Catalogue of Cell Lines and Hybridomas (7th edition, (1992)).

Appropriate vectors for expressing the proteins of the invention in insect cells are usually derived from the SF9 baculovirus. Suitable insect cell lines include mosquito larvae, silkworm, armyworm, moth and *Drosophila* cell lines such as a Schneider cell line (*See* Schneider <u>J. Embryol.</u>

30   <u>Exp. Morphol.</u> 27:353-365 (1987).

As indicated above, the vector, *e.g.*, a plasmid, which is used to transform the host cell, preferably contains DNA sequences to initiate transcription and sequences to control the translation of the protein. These sequences are referred to as expression control sequences.

As with yeast, when higher animal host cells are employed, polyadenylation or

35   transcription terminator sequences from known mammalian genes need to be incorporated into the vector. An example of a terminator sequence is the polyadenylation sequence from the bovine growth hormone gene. Sequences for accurate splicing of the transcript may also be included. An example of a splicing sequence is the VP1 intron from SV40 (Sprague, J. *et al.*, <u>J. Virol.</u> 45: 773-781 (1983)).

Additionally, gene sequences to control replication in the host cell may be

40   incorporated into the vector such as those found in bovine papilloma virus type-vectors.

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Saveria-Campo, M., 1985, "Bovine Papilloma virus DNA a Eukaryotic Cloning Vector" in DNA Cloning Vol. II a Practical Approach Ed. D.M. Glover, IRL Press, Arlington, Virginia pp. 213-238.

The host cells are competent or rendered competent for transformation by various means. There are several well-known methods of introducing DNA into animal cells. These include: calcium phosphate precipitation, fusion of the recipient cells with bacterial protoplasts containing the DNA, treatment of the recipient cells with liposomes containing the DNA, DEAE dextran, electroporation and micro-injection of the DNA directly into the cells.

The transformed cells are cultured by means well known in the art (Biochemical Methods in Cell Culture and Virology, Kuchler, R.J., Dowden, Hutchinson and Ross, Inc., (1977)). The expressed polypeptides are isolated from cells grown as suspensions or as monolayers. The latter are recovered by well known mechanical, chemical or enzymatic means.

2.   Purification

The proteins produced by recombinant DNA technology may be purified by standard techniques well known to those of skill in the art. Recombinantly produced proteins can be directly expressed or expressed as a fusion protein. The protein is then purified by a combination of cell lysis (e.g., sonication) and affinity chromatography. For fusion products, subsequent digestion of the fusion protein with an appropriate proteolytic enzyme releases the desired polypeptide.

The polypeptides of this invention may be purified to substantial purity by standard techniques well known in the art, including selective precipitation with such substances as ammonium sulfate, column chromatography, immunopurification methods, and others. See, for instance, R. Scopes, Protein Purification: Principles and Practice, Springer-Verlag: New York (1982), incorporated herein by reference. For example, in an embodiment, antibodies may be raised to the proteins of the invention as described herein. Cell membranes are isolated from a cell line expressing the recombinant protein, the protein is extracted from the membranes and immunoprecipitated. The proteins may then be further purified by standard protein chemistry techniques as described above.

3.   Antibodies

As mentioned above, antibodies can also be used for the screening of polypeptide products encoded by the polymorphic nucleic acids of the invention. In addition, antibodies are useful in a variety of other contexts in accordance with the present invention. Such antibodies can be utilized for the diagnosis of HH and, in certain applications, targeting of affected tissues.

Thus, in accordance with another aspect of the present invention a kit is provided that is suitable for use in screening and assaying for the presence of polypeptide products encoded by the polymorphic nucleic acids of the invention by an immunoassay through use of an antibody which specifically binds to polypeptide products encoded by the polymorphic nucleic acids of the invention in combination with a reagent for detecting the binding of the antibody to the gene product.

Once hybridoma cell lines are prepared, monoclonal antibodies can be made through conventional techniques of priming mice with pristane and interperitoneally injecting such mice with the hybrid cells to enable harvesting of the monoclonal antibodies from ascites fluid.

In connection with synthetic and semi-synthetic antibodies, such terms are intended to cover antibody fragments, isotype switched antibodies, humanized antibodies (mouse-human, human-

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mouse, and the like), hybrids, antibodies having plural specificities, fully synthetic antibody-like molecules, and the like.

This invention also embraces diagnostic kits for detecting DNA or RNA comprising a polymorphism of Table 1 in tissue or blood samples which comprise nucleic acid probes as described herein and instructional material. The kit may also contain additional components such as labeled compounds, as described herein, for identification of duplexed nucleic acids.

The following examples are provided to illustrate the invention but not to limit its scope. Other variants of the invention will be readily apparent to one of ordinary skill in the art and are encompassed by the appended claims.

F. EXPERIMENTAL EXAMPLES

1. Megabase transcript map

In these studies direct selection, exon-trapping, and genomic sample sequencing were used to generate a transcript map of a 1 megabase region approximately 8.5 megabases telomeric to HLA-A in the vicinity of HFE. This region 6p21.3 was flanked by the genetic markers D6S2242 and D6S2241. The starting material for these experiments was a 1 megabase YAC labeled y899g1 and a bacterial clone contig of this region (Feder et al. Nature Genetics 13:399-408 (1996)). These techniques and other methods used in the study are outlined below.

a. Direct Selection (DS)

Poly A$^+$ RNA from human fetal brain, liver and small intestine (Clontech, Palo Alto, CA) were converted into cDNA using random primers and a Superscript cDNA synthesis kit (Life Technologies, Gaithersburg, MD). The cDNA was digested with Mbo I and ligated to cDNA Mbo I linker-adaptors. Unligated linker-adaptor were removed by passage through cDNA spun columns (Pharmacia, Piscataway, NJ). The 5 ng of each of the ligated cDNAs were amplified using the cDNA Mbo I-S primer (5'-CCTGATGCTCGAGTGAATTC-3'). The amplified products were purified on S-400 spin columns (Pharmacia, Piscataway, NJ), ethanol precipitated and resuspended at 1mg/ml in TE. Gel-purified yac899g1 (Centre d'Etude du Polymorphisme Humain) was processed as described by Morgan et al. (Nucl. Acids Res. 20:5173-5179 (1992)). The cDNAs were mixed in equal molar amounts for a total of 3 mg, and blocked with a mixture of 4 mg Cot-1 DNA (Life Technologies, Gaithersburg, MD), and a cocktail of Sau 3A-digested ribosomal and five different histone DNAs. The blocked cDNAs were hybridized to biotinylated yac899g1 DNA and streptavidin capture was carried out as described by Morgan et al. (ibid). After the second round of selection, the eluted cDNAs were amplified using the cDNA Mbo I-S primer which included a (CUA)4 repeat at the 5' end to facilitate cloning into a version of pSP72 (Promega, Madison, WI) constructed for use with uracil-DNA glycolyase cloning (UDG, Life Technologies, Gaithersburg, MD). Recombinants were transformed in DH5α, 1000 clones picked into a 96 well format, and clones prepped for DNA sequencing using AGTC boiling 96-well mini-prep system (Advance Genetic Technologies, Gaitherburg, MD).

Four hundred and sixty five clones were sequenced and the resulting data searched by BLAST (Altschul et al. J. Mol. Biol. 215:403-410 (1990)). Those clones representing repetitive, bact rial, yeast, mitochondrial and histone sequences wer eliminated from future considerations. The remaining sequenc s were then searched for overlaps and assembled into 108 unique DS contigs.

25

The number of clones per DS contig varied between 1 to 22 with the length of each contig ranging from 250bp to 850 bp. Small sequence-tag-sites PCR assays were developed for each DS contig and two experiments were carried out concomitantly; mapping each DS contig back to the bacterial clone contig of the region and testing for the presence of each DS contig in cDNA libraries. Overall, 86 or 80% of the DS contigs mapped back to the region and were found to be in cDNA libraries. The number of 80% mapping to the region was probably an underestimate of the fidelity of the direct-selection since PCR assays which cross exon-intron boundaries would be expected to fail or give larger size products, thereby being scored negative.

b. Exon-Trapping

CsCl-purified genomic P1 (Genome Systems), BAC (Research Genetics) and PAC (Genome Systems) DNAs were digested with BamHI, Bgl II, Pst I Sac 1 and Xho I and 125 ng of each digest ligated into 500 ng pSPL3 (Church *et al.* Nature Genetics 6:98-105 (1994)) (Life Technologies, Gaithersburg, MD) digested with the appropriate restriction enzyme and phosphatased with calf intestinal alkaline phosphatase (USB, Cleveland, OH). One tenth of the ligation was used to transform XL1-Blue MRF' cells (Stratagene, La Jolla, CA) by electroporation. Nine tenths of the electroporation was used to inoculate 10 ml of LB + 100µg/ml of carbenicillen and after overnight growth, DNA was prepared using Qiagen Q-20 tips (Qiagen GmbH, Hilden Germany). The remaining one tenth was plated on LB +100 µg/ml carbenicillen plates to evaluated the efficiency on cloning and to test individual clones for the present of single inserts. COS-7 cells were seed overnight at a density of 1.4 x10^5/well in 6 well dishes. One µg of DNA was transfected using 6ml of Lipofect-Ace. Cytoplasmic RNA was isolated 48 hr post-transfection. RT-PCR was carried out as described by Church *et al.* (ibid) using commercially available reagents Life Technologies, Gaithersburg, MD). The resulting CUA-tailed PCR fragments for each restriction digested bacterial clone were pooled and UDG cloned into pSP72-U (a derivative of pSP72). The DNA was transformed in DH5α and the cells plated onto nylon membranes. After overnight growth, duplicates were made and the DNA hybridized to ^{32}P end-labeled oligos designed to detect various background products associated with the pSPL3 vector. One set of filters was hybridized with the following gel-purified oligos in 6X SSC aqueous hybridization solution at 42° C:

vector-vector splicing	5'-CGACCCAGCAACCTGGAGAT-3'
cryptic donor-1021	5'-AGCTCGAGCGGCCGCTGCAG-3'
cryptic donor-1134	5'-AGACCCCAACCCACAAGAAG-3'

The filters were washed twice in 6X SSC, 10 mM sodium pyrophosphate (NaPPi) at 60°C, 30 mins.

After overnight autoradiography, non-hybridizing clones were picked and grown in 250 µl of LB + 100µg/ml of carbenicillin in 96 well mini-rack tubes. The samples were analyzed by PCR using the secondary PCR primers supplied in the kit (Life Technologies, Gaithersburg, MD) and those clones with inserts greater than 200 bp were selected for sequencing.

Ninety-six exon traps per bacterial clone were sequenced for a total of 768 reactions and the resulting data analyzed by BLAST. In addition, each potential exon was searched against a database of the 86 DS contigs to eliminate redundant sequences. PCR assays were developed for

each of the potential exons and they were tested for their presence in cDNA libraries. A total of 48 potential exons remained after these screening steps.

c. Sample Sequencing

A minimal set of bacterial clones chosen to cover y899g1 were prepped with the Qiagen Maxi-Prep system and purified on CsCl. Ten micrograms of DNA from each bacterial clone was sonicated in a Heat Systems Sonicator XL and end-repaired with Klenow (USB) and T4 polymerase (USB). The sheared fragments were size selected between three to four kilobases on a 0.7% agarose gel and then ligated to BstXI linkers (Invitrogen). The ligations were gel purified on a 0.7% agarose gel and cloned into a pSP72 derivative plasmid vector. The resulting plasmids were transformed into electrocompetent DH5α cells and plated on LB-carbenicillin plates. A sufficient number of colonies was picked to achieve 15-fold clone coverage. The appropriate number of colonies was calculated by the following equation to generate a single-fold sequence coverage: Number of colonies = size of bacterial clone (in kb)/average sequence read length (0.4 kb). These colonies were prepped in the 96-well AGCT system and end-sequenced with oligo MAP1 using standard ABI Dye Terminator protocols. MAP1 was CGTTAGAACGCGGCTACAAT. The MAP1 sequences were screened locally with the BLAST algorithm against all available public databases. All sequence identities were catalogued and cross referenced to the DS and exon-trapped databases.

A total of 3794 end sequence reactions were run to achieve the theoretical 1X coverage. Eighty-five percent of these sequences contained non-bacterial non-vector inserts. An additional 1060 end sequence reactions were run from the opposite end of the cloning vector to augment the sequence coverage and to prepare for contigging across selected regions. BLAST searches to all publicly available databases identified 12 histone genes and 74 unique expressed sequence fragments (ESF). The ESF represent a collection of ESTs and other expressed sequence fragments that were selected due to their sequence identity over a significant portion of genomic DNA. The ESF were cross referenced against the DS and exon-trapped databases to eliminate redundancies. 58 unique ESF remained, representing 39 distinct clones. Included in these ESF are 5 sequences homologous to histone genes.

Table 3. EST's found by Sample Sequencing Large Insert Bacterial Clones

Clone name	Bacterial clone	Homology 5' blastx	Homology 3' blastx	Poly A+ signal[1]	Genomic poly (A)$_{08}$	cDNA Homology
EST03556	pc157c3	na[2]	none[3]	+	-	cDNA 28
ym33f11	pc157c3	ZNF	na	na	na	
EST04698	pc157c3	na	NSH[4]	+	-	
EST04812	pc157c3	na	NSH	-	-	
yb89b08	pc157c3	NSH	na	na	na	
yd88g11	pc157c3	na	nsh	+	-	
yj49b01	pc157c3	NSH	na	na	na	
yv81d05	pc157c3	HG17 Human	NSH	+	-	cDNA 30
yg57h09	p196e20	BUTYBOVIN	NSH	+	-	cDNA 21
yq23d08	p196e20	BUTYBOVIN	NSH	+	-	cDNA 21

Clone name	Bacterial clone	Homology 5' blastx	Homology 3' blastx	Poly A+ signal[1]	Genomic poly (A)[n,s]	cDNA Homology
yo65f06	p196e20	NSH	na	na	na	cDNA 29
yv88c09	p196e20	BUTYBOVIN	na	na	na	cDNA 29
yd17d06	p196e20	NSH	na	na	na	cDNA 23
ye25g03	p196e20	BUTYBOVIN	NSH	na	na	cDNA 44
ys04h08	pc45p21	NSH	NSH	+	-	cDNA 44
yn01c05	p196e20	BUTYBOVIN	na	na	na	cDNA 32
YG78F10	PC45P21	NSH	NSH	na	na	
yh54f11	p196e20	none	NSH	-	-	
ys05b08	pc157c3	NSH	Alu	-	+	
yb12h11	b132a12	NSH	Histone H3.1	-	-	
HSC2EE082	b132a12	na	NSH	+	-	
HUM160h11b	b132a12	none	na	na	na	
yg04f09	b132b12	Line element	Alu	-	+	
yd37d11	b132a12	NSH	Alu	-	+	
ym29g03	b132a12	Histone H2A	NSH	+	-	cDNA 37
yi77b02	b132a12	NSH	NSH	-	-	cDNA 37
yh76b05	b132a12	NSH	Alu	-	-	
yu98e02	b132a12	NSH	Alue	-	+	
yd72h12	b132a12	Alu	NSH	+	+	
yd19d03	pc222k22	Histone H2B.1	NSH	+	-	
ye98g01	b132a12	NSH	NSH	+	-	cDNA
yi61f07	b132a12	NSH	NSH	-	+	
ESTO5340	b3e17	na	Alu	-	+	
yd35d05	pc222k22	NSH	NSH	-	+	
yc52a05	pc75L14	NSH	na	na	na	
yd84a05	pc75L14	none	none	-	?[5]	
yr42a05	pc75L14	NaPi transport	none	+	-	cDNA 22B
yd83h08	b20h20	NSH	none	+	-	
ye38c09	b20h20	NSH	Alu	-	+	
yp74c05	b20h20	NaPi transport	Alu	?[6]	na	

Bracketed area is the critical region

1	Signal of ATAAA or ATTAA	4	No Significant Homologies
2	Not available	5	3' splice that is not on contig
3	"NONE" reported by blast	6	Poor EST sequence

d. cDNA library screening

Superscript plasmid cDNA libraries, brain, liver and testis, were purchased from Life Technologies, Gaithersburg, MD. Colonies were plated on Hybond N filters (Amersham) using

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standard techniques. Insert probes from DS, exons and EST (I.M.A.G.E. clones; Genome Systems) were all isolated by PCR followed by purification in low-melting point agarose gels (Seakem). The DNAs were labeled in gel using the Prime-it II kit (Stratagene, La Jolla, CA). Small exon probes were labeled using their respective STS PCR primers instead of random primers. Up to 5 different probes

5   were pooled in a hybridization. Filters were hybridized in duplicate using standard techniques. Putative positives were screened by PCR using the probe's STSs to identify clones. Inserts from positive clones were subcloned in pSP72 and sequenced.

e. Northern blots and RT-PCR analysis

Multiple tissue northern blots were purchased from Clontech and hybridized according

10   the manufacturer's instructions. RT-PCR was carried out on random primed first strand cDNA made from poly A+ RNA (Clontech) using AmpliTaq Gold (Perkin-Elmer). Control reactions were performed on RNA samples processed in the absence of reverse transcriptase to control for genomic DNA contamination.

f. Genomic Sequencing

15   The MAP1 sequences from the bacterial clones b132a2, 222K22, and 75L14 were assembled into contigs with the Staden package (available from Roger Staden, MRC). A minimal set of 3 kb clones was selected for sequencing with oligo labeled MAP2 that sits on the opposite end of the plasmid vector. The sequence of MAP2 was GCCGATTCATTAATGCAGGT. The MAP2 sequences were entered into the Staden database in conjunction with the MAP1 sequences to generate a tiling

20   path of 3 kb clones across the region. These sequences were also screened with the BLAST algorithm and all novel sequence identities were noted. The plasmid 3 kb libraries were concurrently transformed in 96 well format into pox38UR (available from C. Martin, Lawrence Berkeley Laboratories). The transformants were subsequently mated with JGM (Strathman et al. P.N.A.S. 88:1247-1250 (1991) in 96 well format. All matings of the 3 kb clones within the tiling path were

25   streaked on LB-carbenicillin-kanamycin plates and a random selection of 12 colonies per 3 kb clone was prepped in the AGCT system. The oligos -21: CTGTAAAACGACGGCCAGTC, and REV: GCAGGAAACAGCTATGACC were used to sequence off both ends of the transposon. Each 3 kb clone was assembled in conjunction with the end sequence information from all bacterial clones to generate complete sequence across the region. The genomic sequence was analyzed with the

30   BLAST nucleotide and protein homology algorithms and the GRAIL 1.2 software to identify novel open reading frames (ORF) for gene finding.

g. Discussion

A compilation of 174 ESF led to the construction of an expressed sequence map of the region that served as the framework for the isolation of full-length cDNAs (Figure 1). (The map

35   shows the subset of ESF that were actually mapped). Probes were developed for 82 best ESFs which appeared to be derived from the coding portions of cDNAs and the appropriate cDNA libraries were screened. This led to the isolation of 19 cDNAs, 17 of which represented novel sequences. 70 of the 174 ESF were included in the cDNAs isolated (40%). 36 probes failed to produce any clones even after repeated screening of several libraries. 51 ESF which were not accounted for in the cDNAs

cloned were not used in any screen. Therefore, it is possible that some additional genes within this 1 megabase region may have escaped detection.

A list of these cDNAs cloned and a comparison of the methods used to find them is presented in Table 4. Direct selection found 14 out of the 18 cDNAs contained within the boundaries of the YAC used in the experiment. Exon trapping found 15 out of the 19 cDNAs contained within the boundaries of the large insert bacterial clone contig. Sample sequencing identified 11 genes that had corresponding ESTs in the public database.

Table 4. Comparison of gene finding methods

Bacterial Clone	CDNA #	Homology	EST	DS	Exon Trap
157c	28	zinc finger	EST03556	2	1
157c3	30	nonhistone	yv81d05	1	none
			yvh07a10		
157c3	46	ORF	yd88g11	1	
157c3	20	BT	none	none	3
p18696	21	BTF1	yn01G5	4	5
			yg23d08		
			yg57h09		
			yu15h03		
45p21	32	BTF2	yg78f10	7	3
			yn01c05		
45p21	29	BTF3	ye25g03	2	9
			yo65f06		
45p21	23	BTF4	yd17d06	4	6
45p21	44	BTF5	ys04h08	2	4
3e17	41	genomic?	none	none	1
132a2	43	genomic?	none	none	3
132a2	36	genomic?	none	1	none
132a2	37	histone 2A	ym29g03	3	none
			yh87a03		
75114	24	MHC class 1	ye98g01	1	2
132a2	39	genomic?	none	none	4
132a2	27	Ro/SSA	none	3	4
132a2	22B	NPT1-like	yr42a05	1	7
			yf09g06		
20h20	22E	NPT1-like	none	2	5
20h20	NPT1	NPT1	yp74c05	N/A	3

30

As a final approach, a tiling path with overlapping end sequences from the sample sequence database was generated. Each 3 kb clone within the path was shotgun-sequenced using transposable elements as platforms for dual end sequencing. These individual clones were assembled in conjunction with the end sequences from all bacterial clones in the region. The resulting sequence

5   (Figure 2) was analyzed systematically with BLAST homology searches and the Grail 1.2 program to identify novel open reading frames (ORF) and other gene-like structures. The BLAST homology searches did not produce any probes that had not already been identified by sample sequencing. Grail predicted exons for all the genes in the region, but was only able assemble the histones into any representative form. A detailed analysis of BLAST homology searches to protein databases identified

10   an enticing homology to a zinc alpha 2 glycoprotein approximately 25 kb upstream of HFE, but the lack of a substantial ORF and the presence of a stop codon suggested that it was a pseudogene. Figure 2 shows the positions, the exon and intron structures, and the relative orientation of transcription of novel genes within this region. Also shown are the positions and transcriptional orientations of the histone genes. A total of 12 histone genes were identified in this study.

15   In an effort to account for the ESTs that did not associate with the characterized genes in the 250 kb region, the genomic sequence around the putative 3' ends were examined for polyadenylation signals to determine whether certain EST sequences may have originated from genomic DNA contamination in the normalized cDNA libraries used in EST generation. The positions of the 14 ESTs found in this region are indicated in Figure 2 to show those associated with the cDNAs

20   cloned and those which did not associate with genomic DNA of obvious coding potential. Four ESTs corresponded to 3 of the 4 cDNAs cloned from the region (Table 2). One EST encoded a histone H2B.1 gene and another was a repetitive element. Of the remaining 8, 6 EST clones were used as probes of cDNA libraries with negative results. Those sequences representing putative 3' ends of cDNA were searched for the presence of poly (A)+ addition signals. Five of the 13 ESTs which had 3'

25   end sequence, had the sequence ATAAA or ATTAA. Five of the remaining 8 ESTs that did not have a poly (A)+ addition signal had genomic encoded stretches of poly (A) near the end of EST sequence and, therefore, may have been created by oligo d(T) priming of contaminating genomic DNA. This analysis was expanded to include all ESTs in the large-insert bacterial contigs with definitive 3' ends. Of the remaining 26, 15 had 3' end sequence and, of these, 8 had poly (A)+ addition signals. Five of

30   these 8 ESTs were associated with the cloned cDNAs. Of the remaining 7 which did not have poly (A)+ addition signals, 4 had genomic encoded stretches of poly (A).

i. Butyrophilin gene family

The human homolog of the bovine butyrophilin gene (BT) was cloned and mapped to approximately 480 kb centromeric to HFE (Figure 1). BT is a transmembrane protein of unknown

35   function which constitutes 40% of the total protein associated with the fat globule of bovine milk (Jack et al. J. Biol. Chem. 265:14481-14486 (1990)). A human homolog of BT has recently been cloned by Tayloer et al. (Biochem Biophys Acta 1306:1-4 (1996)). The results in this study indicated that BT is a member of a gene family with at least five other members of the family residing in this region (Figure 1). A comparison of these proteins is shown in Figure 3. The proteins were aligned based on their

40   descending order of relatedness and to minimized gaps in the sequence. Each of the five proteins

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display varying degrees of homology to BT. BTF1 (cDNA 21), BTF2 (cDNA 32), BTF5 (cDNA 44), and BTF3 (cDNA 29) are 45%, 48%, 46%, and 49%, identical to BT, whereas BTF4 (cDNA 23), which is more similar to BTF3 (cDNA 29), is only 26% identical. This low degree of identity to BT is largely due to a truncation at the carboxyl terminus of the protein. The BTF family falls into two groups: BTF1 and 2 which are more related to each other than to BT or the other BTF members, and BTF5, 3 and 4, which appear to have a common evolutionary origin. The order of these genes on the chromosome suggests that the BT gene has duplicated two times, giving rise to BTF1 and BTF5. Subsequently, it appears likely these two genes experienced further duplication events to give rise to the other members in their groups.

The three major components of BT, the B-G immunoglobulin superfamily domain (containing the V consensus sequence) (Miller *et al.* Proc. Natl. Acad. Sci. U.S.A. 88:4377-4381 (1991)), the transmembrane region, and the B30-2 exon are found in all of these proteins (with the exception of BTF4 (cDNA 23) which lacks the B30-2 exon by virtue of the carboxyl terminal truncation). The exon B30-2 is a previously noted feature of the MHC class 1 region found approximately 200 kb centromeric to the HLA-A gene (Vernet *et al.*, J. Mol. Evol. 37:600-612 (1993)). In addition this exon is found in several genes of diverse function telomeric to HLA-A namely MOG (approximately 200 kb) and RFP (approximately 1 megabase) (Amadou *et al.* Genomics 26:9-20 (1995)).

The levels of the BTF mRNA were analyzed by northern blot analysis (Figure 4A). The expression of the BTF genes fell into two patterns. BTF1 and BTF2 were expressed as a single major transcript of 2.9 kb and one minor transcript of 5.0 kb. These genes were expressed at high levels in all the tissues tested with the exception of the kidney where the expression level was less. The two genes are 90% identical at the DNA sequence level, therefore, it is possible that the signal observed on the northerns was the result of cross-hybridization and only one of the two genes was actually expressed. To address this possibility RT-PCR experiments were carried out on a panel of different tissues in order to detect possible tissue dependent expression that would suggest that both genes are expressed. Identical, and thus equivocal, results were obtained with both BTF1 and BTF2 amplification (Figure 4B).

The second group of genes, BTF3-5, are expressed as three (BTF5) (Figure 4A) and two (BTF3 and 4) transcripts ranging from 4.0 to 3.3 kb. BTF5 is expressed at moderate levels in all tissues tested with the exception of the kidney where the expression level is less. RT-PCR experiments showed that mRNA from the BTF5 gene can be found in all tissues tested, including the kidney (Figure 4B). Identical results were obtained with primers from the other genes of this group (data not shown). These genes are also 90% identical to each other at the DNA sequence level (but only 58% identical to BTF1 and 2), hence like BTF1 and BTF2, cross-hybridization could account for the similarity in size and patterns on the northern blots and RT-PCR. This might be particularly true for BTF4 which lacks the B30-2 exon but still hybridizes to larger size transcripts like BTF5 and BTF3.

ii. A gene with similarity to 52 kD Ro/SSA auto-antigen

Located approximately 120 kb telomeric to the HFE gene is a gene, RoRet, that has 58% amino acid similarity to the 52 kD Ro/SSA protein, an auto-antigen of unknown function that is frequently recognized by antibodies in patients with systemic lupus and Sjogren's syndrome (Anderson

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et al. Lancet 2:456-560 (1961); Clark et al. J. Immunol. 102:117-122 (1969)) (Figures 1 and 2).

Alignment of the predicted amino acid sequence of this cDNA with that of 52 kD Ro/SSA indicated two features associated with the 52 kD Ro/SSA protein: a putative DNA binding cysteine rich motif (C-X-(I,V)-C-X(11-30)-C-X-H-X-(F,I,L)-C-X(2)-C-(I,L,M)-X(10-18)-C-P-X-C) found at the N terminus

5   (Freemont et al. Cell 64: 483-484 (1991)) and the B30-2 exon found near the carboxyl terminus, are both conserved in RoRet (Figure 5). Northern blot analysis indicated the RoRet gene was expressed as two major transcripts of 2.8 and 2.2 kb and two minor transcripts of 7.1 and 4.4 kb in all of the tissues on the blot at levels reflective of the RNA amounts as determined by β-actin probing (Figure 6A). Using RT-PCR, expression can also be detected in small intestine, kidney liver, and spleen

10   (Figure 6B).

iii. Two genes with homology to a sodium phosphate transporter

A cDNA for a sodium phosphate transport protein (NPT1) was previously cloned and mapped to 6p21.3 using a somatic cell hybrid panel (Chong et al. Genomics 18:355-359 (1993)). NPT1 maps 320 kb telomeric to the HFE gene (Figures 1 and 2). Two additional cDNAs were cloned

15   which show appreciable homology to NPT1 (Figure 5). These genes, NPT3 and NPT4, mapped 1.5 megabases and 1.3 megabases centromeric to the NPT1 gene (Figure 1). Like NPT1, the gene products of NPT3 and NPT4 were extremely hydrophobic, which may reflect a membrane location. Both proteins gave hydrophilicity profiles which were indistinguishable from NPT1 in this study (data not shown). Northern blot analysis indicated that the two genes have different patterns of expression

20   (Figure 6C). NPT3 was expressed at high levels as a 7.2 kb transcript predominately in muscle and heart. Lesser amount of the mRNA were also found in brain, placenta, lung, liver and pancreas. RT-PCR analysis indicated that expression of the proper size PCR fragment for NPT3 was clearly absent in fetal brain, bone marrow and small intestine (Figure 6D). A smaller size fragment was detectable in all tissues with the exception of the liver, which may represent evidence for alternative

25   splicing. Although expression was apparently absent from the kidney by northern blot analysis, it was detectable by RT-PCR. Expression was also noted in the mammary gland, spleen and testis. NPT4, on the other hand, was expressed only in the liver and the kidney as a smear of transcripts approximately 2.6 - 1.7 kb (Figure 6C). RT-PCR confirmed these results, although a small amount of the proper size PCR fragment was also found in the small intestine and testis (Figure 6D). Other

30   tissues showed amplification, but the fragments were of larger and smaller size than that produced by the cDNA 22E positive control. Hence, these two genes which apparently have the structural characteristics of a sodium phosphate transporter, appeared to be under the control of different regulatory mechanism that lead to differential patterns of expression.

2.   Sequencing of 235 kb from a Homozygous Ancestral (Affected) Individual

35   In these studies the entire genomic sequence was determined from an HH affected individual for a region corresponding to a 235,033 bp region surrounding the HFE gene between the flanking markers D6S2238 and D6S2241. The sequence was derived from a human lymphoblastoid cell line, HC14, that is homozygous for the ancestral HH mutation and region. The sequence from the ancestral chromosome (Figure 9) was compared to the sequence of the region in an unaffected

40   individual (Figure 8) disclosed in copending U.S.S.N. 08/724,394 to identify polymorphic sites. A

subset of the polymorphic alleles so defined were further studied to determine their frequency in a collection of random individuals.

The cell line HC14 was deposited with the ATCC om June 25, 1997, and is designated ATCC CRL-12371.

5 ### a. Cosmid Library Screening

The strategy and methodology for sequencing the genomic DNA for the affected individual was essentially as described in copending U.S.S.N. 08/724,394, hereby incorporated by reference in its entirety. Basically, a cosmid library was constructed using high molecular weight DNA from HC14 cells. The library was constructed in the supercos vector (Stratagene, La Jolla, CA).

10 Colonies were replicated onto Biotrans nylon filters (ICN) using standard techniques. Probes from genomic subclones used in the generation of the sequence of the unaffected sequence disclosed in 08/724,394 were isolated by gel electrophoresis and electroporation. Subclones were chosen at a spacing of approximately 20 kb throughout the 235 kb region. The DNA was labeled by incorporation of 32P dCTP by the random primer labeling approach. Positively hybridizing clones were isolated to

15 purity by a secondary screening step. Cosmid insert ends were sequenced to determine whether full coverage had been obtained, and which clones formed a minimal path of cosmids through the 235 kb region.

b. Sample Sequencing

A minimal set of cosmid clones chosen to cover the 235 kb region were prepped with

20 the Qiagen Maxi-Prep system. Ten micrograms of DNA from each cosmid preparation were sonicated in a Heat Systems Sonicator XL and end-repaired with Klenow (USB) and T4 DNA polymerase (USB). The sheared fragments were size selected between three to four kilobases on a 0.7% agarose gel and then ligated to BstXI linkers (Invitrogen). The ligations were gel purified on a 0.7% agarose gel and cloned into a pSP72 derivative plasmid vector. The resulting plasmids were transformed into

25 electrocompetent DH5α cells and plated on LB-carbenicillin plates. A sufficient number of colonies was picked to achieve 15-fold clone coverage. The appropriate number of colonies was calculated by the following equation to generate a single-fold sequence coverage: Number of colonies = size of bacterial clone (in kb)/average sequence read length (0.4 kb). These colonies were prepped in the 96-well Qiagen REAL, and the 5' to 3' DNA Prep Kit, and AGCT end-sequenced with oligo MAP1 using

30 standard ABI Dye Terminator protocols. MAP1 was CGTTAGAACGCGGCTACAAT.

c. Genomic Sequencing

The MAP1 sequences from the cosmid clones HC182, HC187, HC189, HC195, HC199, HC200, HC201, HC206, HC207, and HC212 were assembled into contigs with the Staden package (available from Roger Staden, MRC). A minimal set of 3 kb clones was selected for

35 sequencing with oligo labeled MAP2 that sits on the opposite end of the plasmid vector. The sequence of MAP2 was GCCGATTCATTAATGCAGGT. The MAP2 sequences were ent red into the Staden database in conjunction with the MAP1 sequences to generate a tiling path of 3 kb clones across the region. The plasmid 3 kb libraries were concurrently transformed in 96 well format into pox38UR (available from C. Martin, Lawrence Berkeley Laboratories). The transformants were subsequently

40 mated with JGM (Strathman et al. P.N.A.S. 88:1247-1250 (1991) in 96 well format. All matings of the

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3 kb clones within the tiling path were streaked on LB-carbenicillin-kanamycin plates and a random selection of 12 colonies per 3 kb clone was prepped in the AGCT system. The oligos -21: CTGTAAAACGACGGCCAGTC, and REV: GCAGGAAACAGCTATGACC were used to sequence off both ends of the transposon. Each 3 kb clone was assembled in conjunction with the end sequence information from all cosmid clones in the region.

In some regions, the coverage of the genomic sequence by cosmids was incomplete. Any gaps in the sequence were filled by using standard PCR techniques to amplify genomic DNA in those regions and standard ABI dye terminator chemistry to sequence the amplification products.

d. Identification of Polymorphic Sites

The assembled sequence of the cosmid clones in connection with the PCR amplified genomic DNA was compared to the genomic sequence of the unaffected individual using the FASTA algorithm. Numeric values were assigned to the sequenced regions of 1 to 235,303, wherein base 1 refers to the first C in the CA repeat of D6S2238 and base 235,303 is the last T in the GT repeat of D6S2241 of the underlined unaffected sequence (Figure 8). Table 1 lists the differences between the two compared sequences. Note that previously disclosed (Feder et al., Nature Genetics 13:399-408 (1996)) polymorphic sites D6S2238 (base 1), D6S2241 (base 235,032), 24d1 (base 41316), and D6S2239 (base 84841) are not included in the list of new polymorphisms, although they are provided for reference in a footnote to the Table and were observed in the ancestral sequence. In the Table, a single base change such as C-T refers to a C in the unaffected sequence at the indicated base position that occurred as a T in the corresponding position in the affected sequence. Similarly, an insertion of one or more bases, such as TTT in the affected sequence, is represented as "TTT INS" between the indicated bases of the unaffected sequence. A deletion of one or more bases occurring in the affected sequence, such as AAA DEL, is represented as the deletion of the indicated bases in the unaffected sequence.

e.   Characterization of Rare Polymorphisms

In this study about 100 of the polymorphisms of Table 1 were arbitrarily chosen for further characterization. Allele frequencies in the general population were estimated by OLA analysis using a population of random DNAs (the "CEPH" collection, J. Dausset et al., Genomics 6(3):575-577 (1990)). These results are provided in Table 2.

One single base pair difference, occurring at base 35983 and designated C182.1G7T/C (an A to G change on the opposite strand) was present in the ancestral chromosome and rare in the random DNAs. This change occurred in a noncoding region of the hemochromatosis gene near exon 7 approximately 5.3 kb from the 24d1 (Cys282Tyr) mutation. OLA was used to genotype 90 hemochromatosis patients for the C182.1G7T/C base pair change. The frequency for C occurring at this position in the patients was 79.4% as compared to 5% in the random DNAs. Eighty-five of the 90 patients assayed contained identical 24d1 and C182.1G7T/C genotypes. Four of the remaining 5 patients were homozygous at 24d1 and heterozygous at C182.1G7T/C; one was heterozygous at 24d1 and homozygous at C182.1G7T/C. The primers used for this analysis were as follows.

PCR primers for detection:

     182.1G7.F   5'-GCATCAGCGATTAACTTCTAC -3'

     182.1G7.R   5'-TTGCATTGTGGTGAAATCAGGG -3'

For the detection assay, the biotinylated primers used were as follows.

     182.1G7.C   5' (b)CTGAGTAATTGTTTAAGGTGC -3'

     182.1G7.T   5' (b)CTGAGTAATTGTTTAAGGTGT -3'

The phosphorylated digoxigenin-labeled primer used was:

     182.1G7.D   5' (p)AGAAGAGATAGATATGGTGG -3'

A further rare single base pair change was detected at 61,465bp. The inheritance pattern of this polymorphism, C195.1H5C/T (a G to A change on the opposite strand), is identical to that of 24d1. The frequency of T occurring at that position (C195.1H5T) observed in a set of 76 patients was 78.5% as compared to 5% in random individuals.

PCR primers for detection:

     1951H5.3F   5'-GAATGTGACCGTCCCATGAG-3'

     1951H5.3R   5'-CAACTGAATATGCAGAAAAAAGTACACC-3'

For the detection assay, the biotinylated primers used were:

     1951H5.3.4   5' (b)AGTAGCTGGGACTCACGGTGT-3'

     1957H5.3.5   5' (b)AGTAGCTGGGACTCACGGTGC-3'

The phosphorylated digoxigenin-labeled primer used was:

     1951H5.3.6   5' (p)GCGCCACCACTCCCAGCTCAT-3'

These rare alleles are thus preferred surrogate markers for 24d1 and are especially useful in screening assays for the likely presence of 24d1 and/or 24d2.

All publications, patents, and patent applications cited herein are hereby incorporated by reference in their entirety.

WHAT IS CLAIMED IS:

1   1.   An oligonucleotide comprising at least 8 to about 100 consecutive bases from the
2   sequence of Figure 9, or the complement of the sequence, wherein the at least 8 to about 100
3   consecutive bases includes at least one polymorphic site of Table 1.

1   2.   The oligonucleotide of claim 1, wherein the polymorphic site is selected from the
2   group consisting of base 35983 or base 61465.

1   3.   An oligonucleotide pair selected from the sequence of Figure 9 or its complement for
2   amplification of a polymorphic site of Table 1.

1   4.   An isolated nucleic acid molecule comprising about 100 consecutive bases to about
2   235 kb substantially identical to the sequence of Figure 9, wherein the DNA molecule comprises at
3   least one polymorphic site of Table 1.

1   5.   The isolated nucleic acid molecule of claim 4, wherein the polymorphic site is selected
2   from the group consisting of base 35983 or base 61465.

1   6.   The isolated nucleic acid molecule of claim 4, wherein the nucleic acid is selected
2   from the group consisting of cDNA, RNA, or genomic DNA.

1   7.   A polypeptide encoded by the nucleic acid molecule of claim 4.

1   8.   An antibody which specifically recognizes the polypeptide of claim 7.

1   9.   A method to determine the presence or absence of the common hereditary
2   hemochromatosis (HFE) gene mutation in an individual comprising:
3   providing DNA or RNA from the individual; and
4   assessing the DNA or RNA for the presence or absence of a haplotype of Table 1,
5   wherein, as a result, the absence of a haplotype of Table 1 indicates the likely absence of the
6   HFE gene mutation in the genome of the individual and the presence of the haplotype indicates the
7   likely presence of the HFE gene mutation in the genome of the individual.

1   10.   The method of claim 9, wherein the method further comprises assessing the RNA or
2   DNA for the presence of at least one of the polymorphisms 24d1, 24d2, HHP-1, HHP-19, or HHP-29;
3   or microsatellite repeat alleles 19D9:205, 18B4:235, 1A2:239, 1E4:271, 24E2:245, 2B8:206, 3321-
4   1:98, 4073-1:182, 4440-1:180, 4440-2:139, 731-1:177, 5091-1:148, 3216-1:221, 4072-2:170, 950-
5   1:142, 950-2:164, 950-3:165, 950-4:128, 950-6:151, 950-8:137, 63-1:151, 63-2:113, 63-3:169, 65-

6   1:206, 65-2:159, 68-1:167, 241-5:108, 241-29:113, 373-8:151, 373-29:113, D6S258:199, D6S265:122,
7   D6S105:124, D6S306:238, D6S464:206, or D6S1001:180.

1   11.   The method of claim 9, wh rein the haplotype comprises at least two polymorphic
2   sites of Table 1.

1   12.   The method of claim 11, wherein one of the at least two polymorphic sites of Table 1
2   is at base 35983 or 61465.

1   13.   The method of claim 11, wherein the haplotype comprises at least three polymorphic
2   sites of Table 1.

1   14.   A method to determine the presence or absence of the common hereditary
2   hemochromatosis (HFE) gene mutation in an individual comprising:
3   providing DNA or RNA from the individual; and
4   assessing the DNA or RNA for the presence or absence of a genotype defined by a
5   polymorphic allele of Table 1,
6   wherein, as a result, the absence of a genotype defined by a polymorphic allele of Table 1
7   indicates the likely absence of the HFE gene mutation in the genome of the individual and the
8   presence of the genotype indicates the likely presence of the HFE gene mutation in the genome of the
9   individual.

1   15.   The method of claim 15, wherein the polymorphic allele occurs in less than about 50%
2   of a random population of individuals.

1   16.   The method of claim 15, wherein the polymorphic allele occurs in less than about 25%
2   of a random population of individuals.

1   17.   The method of claim 15, wherein the polymorphic allele occurs in less than about 5%
2   of a random population of individuals.

1   18.   The method of claim 15, wherein the genotype is C182.1G7C or C195.1H5T.

1   19.   A kit comprising one or more oligonucleotides of claim 1.

1   20.   A kit comprising at least one oligonucleotide pair of claim 3.

1   21.   A culture of lymphoblastoid cells having the designation ATCC CRL-12371.

1   22.   An isolated nucleic acid sequence comprising a sequence substantially identical to

2   BTF1.

1   23.   The isolated nucleic acid sequence of claim 23, wherein the nucleic acid is cDNA.

1   24.   The polypeptide encoded by the isolated nucleic acid sequence of claim 23.

1   25.   A vector comprising the nucleic acid sequence of claim 23.

1   26.   A host cell stably transfected with the nucleic acid sequence of claim 23.

1   27.   An antibody that is specifically immunoreactive with the polypeptide of claim 24.

1   28.   An isolated nucleic acid sequence comprising a sequence substantially identical to

2   BTF2.

1   29.   The isolated nucleic acid sequence of claim 28, wherein the nucleic acid is cDNA.

1   30.   The polypeptide encoded by the isolated nucleic acid sequence of claim 28.

1   31.   A vector comprising the nucleic acid sequence of claim 28.

1   32.   A host cell stably transfected with the nucleic acid sequence of claim 28.

1   33.   An antibody that is specifically immunoreactive with the polypeptide of claim 30.

1   34.   An isolated nucleic acid sequence comprising a sequence substantially identical to

2   BTF3.

1   35.   The isolated nucleic acid sequence of claim 34, wherein the nucleic acid is cDNA.

1   36.   The polypeptide encoded by the isolated nucleic acid sequence of claim 34.

1   37.   A vector comprising the nucleic acid sequence of claim 34.

1   38.   A host cell stably transfected with the nucleic acid sequence of claim 34.

1   39.   An antibody that is specifically immunoreactive with the polypeptide of claim 36.

1
2   BTF4.
40.   An isolated nucleic acid sequence comprising a sequence substantially identical to

1   41.   The isolated nucleic acid sequence of claim 40, wherein the nucleic acid is cDNA.

1   42.   The polypeptide encoded by the isolated nucleic acid sequence of claim 40.

1   43.   A vector comprising the nucleic acid sequence of claim 40.

1   44.   A host cell stably transfected with the nucleic acid sequence of claim 40.

1   45.   An antibody that is specifically immunoreactive with the polypeptide of claim 42.

1
2   BTF5.
46.   An isolated nucleic acid sequence comprising a sequence substantially identical to

1   47.   The isolated nucleic acid sequence of claim 46, wherein the nucleic acid is cDNA.

1   48.   The polypeptide encoded by the isolated nucleic acid sequence of claim 46.

1   49.   A vector comprising the nucleic acid sequence of claim 46.

1   50.   A host cell stably transfected with the nucleic acid sequence of claim 46.

1   51.   An antibody that is specifically immunoreactive with the polypeptide of claim 48.

1
2   NTP-3.
52.   An isolated nucleic acid sequence comprising a sequence substantially identical to

1   53.   The isolated nucleic acid sequence of claim 52, wherein the nucleic acid is cDNA.

1   54.   The polypeptide encoded by the isolated nucleic acid sequence of claim 52.

1   55.   A vector comprising the nucleic acid sequence of claim 52.

1   56.   A host cell stably transfected with the nucleic acid sequence of claim 52.

1   57.   An antibody that is specifically immunoreactiv with the polypeptide of claim 54.

1   58.   An isolated nucleic acid sequence comprising a sequence substantially identical to

2   NTP-4.

1   59.   The isolated nucleic acid sequence of claim 58, wherein the nucleic acid is cDNA.

1   60.   The polypeptide encoded by the isolated nucleic acid sequence of claim 58.

1   61.   A vector comprising the nucleic acid sequence of claim 58.

1   62.   A host cell stably transfected with the nucleic acid sequence of claim 58.

1   63.   An antibody that is specifically immunoreactive with the polypeptide of claim 60.

1   64.   An isolated nucleic acid sequence comprising a sequence substantially identical to

2   RoRet.

1   65.   The isolated nucleic acid sequence of claim 64, wherein the nucleic acid is cDNA.

1   66.   The polypeptide encoded by the isolated nucleic acid sequence of claim 64.

1   67.   A vector comprising the nucleic acid sequence of claim 64.

1   68.   A host cell stably transfected with the nucleic acid sequence of claim 64.

1   69.   An antibody that is specifically immunoreactive with the polypeptide of claim 66.

1   70.   An isolated nucleic acid sequence comprising at least 18 contiguous nucleotides

2   substantially identical to 18 contiguous nucleotides of BTF1.

1   71.   An isolated nucleic acid sequence comprising at least 18 contiguous nucleotides

2   substantially identical to 18 contiguous nucleotides of BTF2.

1   72.   An isolated nucleic acid sequence comprising at least 18 contiguous nucleotides

2   substantially identical to 18 contiguous nucleotides of BTF3.

1   73.   An isolated nucleic acid sequence comprising at least 18 contiguous nucleotides

2   substantially identical to 18 contiguous nucleotides of BTF4.

1   74.   An isolated nucleic acid sequence comprising at least 18 contiguous nucleotides

2   substantially identical to 18 contiguous nucleotides of BTF5.

1   75.   An isolated nucleic acid sequence comprising at least 18 contiguous nucleotides
2   substantially identical to 18 contiguous nucleotides of NPT3.

1   76.   An isolated nucleic acid sequence comprising at least 18 contiguous nucleotides
2   substantially identical to 18 contiguous nucleotides of NPT4.
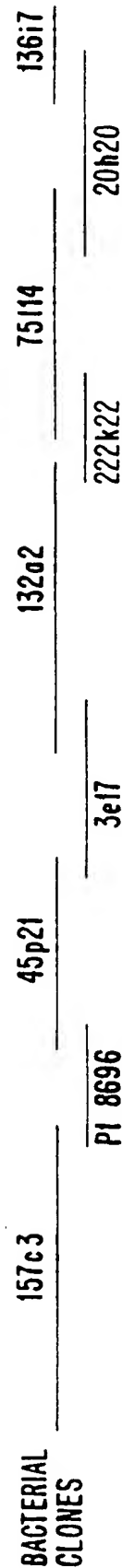
1   77.   An isolated nucleic acid sequence comprising at least 18 contiguous nucleotides
2   substantially identical to 18 contiguous nucleotides of RoRet.
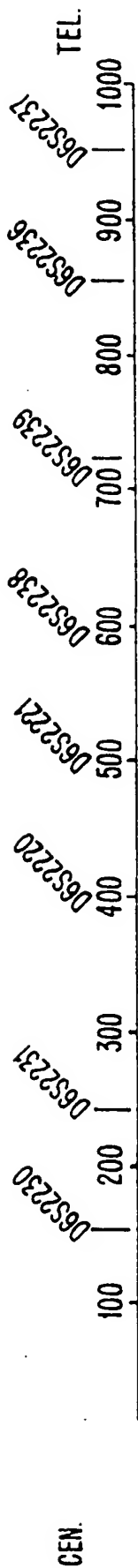
FIG. 1.

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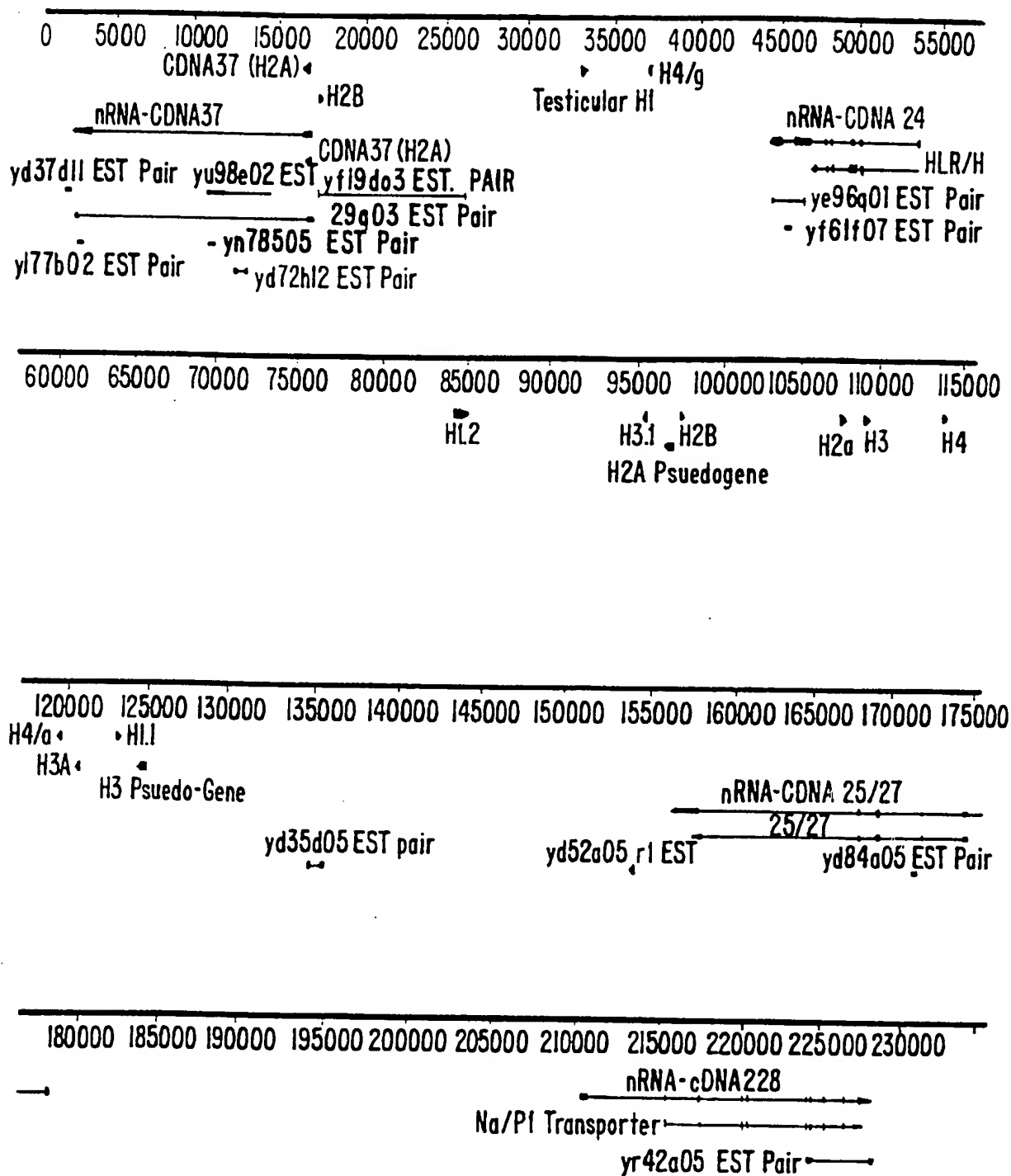FIG. 2.

```
BT      --MAVFPSSGLPRCL---LTLILLQLPKLDSAPFDVIGPPEPILAVVGEDAELPCRLSPN
BTF1    MESAAALHFSRPAS----LLLLLLLSLCALVSAQFIVVGPTDPILATVGENTTLRCHLSPE
BTF2    MEPAAALHFSLPASLLLLLLLLLLSLCALVSAQFTVVGPANPILAMVGENTTLRCHLSPE
BTF5    MKMASFLAFLLLNFR---VCLLLLQLLMPHSAQFSVLGPSGPILAMVGEDADLPCHLFPT
BTF3    MKMASSLAFLLLNFH---VSLFLVQLLTPCSAQFSVLGPSGPILAMVGEDADLPCHLFPT
BTF4    MKMASSLAFLLLNFH---VSLLLVQLLTPCSAQFSVLGPSGPILAMVGEDADLPCHLFPT
               *         .  * *. *    ** *  *.**  **** ***  . *  * . * *


BT      ASAEHLELRWFRKKVSPAVLVHRDGREQEAEQMPEYRGRATLVQDGIAKGRVALRIRGVR
BTF1    KNAEDMEVRWFRSQFSPAVFVYKGGRERTEEQMEEYRGRTTFVSKDISRGSVALVIHNIT
BTF2    KNAEDMEVRWFRSQFSPAVFVYKGGRERTEEQMEEYRGRITFVSKDINRGSVALVIHNVT
BTF5    MSAETMELKWVSSSLRQVVNVYADGKEVEDRQSAPYRGRTSILRDGITAGKAALRIHNVT
BTF3    MSAETMELRWVSSSLRQVVNVYADGKEVEDRQSAPYRGRTSILRDGITAGKAALRIHNVT
BTF4    MSAETMELKWVSSSLRQVVNVYADGKEVEDRQSAPYRGRTSILRDGITAGKAALRIHNVT
              **  .*..*       * *    *.*       *    ****  .    *   *  ** *...


BT      VSDDGEYTCFFREDGSYEEALVHLKVAALGSDPHISMQVQENGEICLECTSVGWYPEPQV
BTF1    AQENGTYRCYFQEGRSYDEAILHLVVAGLGSKPLISMRGHEDGGIRLECISRGWYPKPLT
BTF2    AQENGIYRCYFQEGRSYDEAILRLVVAGLGSKPLIEIKAQEDGSIWLECISGGWYPEPLT
BTF5    ASDSGKYLCYFQDGDFYEKALVELKVAALGSDLHVDVKGYKDGGIHLECRSTGWYPQPQI
BTF3    ASDSGKYLCYFQDGDFYEKALVELKVAALGSDLHIEVKGYEDGGIHLECRSTGWYPQPQI
BTF4    ASDSGKYLCYFQDGDFYEKALVELKVAALGSNLHVEVKGYEDGGIHLECRSTGWYPQPQI
              . *   * * *.*..   *. *.. * ** ***   . ..     * * *** * * **** *


BT      QWRTSKGEKFPSTSESRNPDEEGLFTVAASVIIRDTSTKNVSCYIQNLLLGQEKKVEISI
BTF1    VWRDPYGGVAPALKEVSMPDADGLFMVTTAVIIRDKSVRNMSCSINNTLLGQKKESVIFI
BTF2    VWRDPYGEVVPALKEVSIADADGLFMVTTAVIIRDKYVRNVSCSVNNTLLGQEKETVIFI
BTF5    QWSNNKGENIPTVEAPVVADGVGLYAVAASVIMRGSSGEGVSCTIRSSLLGLEKTASISI
BTF3    KWSDTKGENIPAVEAPVVADGVGLYAVAASVIMRGSSGGGVSCIIRNSLLGLEKTASISI
BTF4    QWSNAKGENIPAVEAPVVADGVGLYEVAASVIMRGGSGEGVSCIIRNSLLGLEKTASISI
              *       *     *.        *   **. *...**.*     ..**  .   *** *    * *


BT      PASSLPRLTPWIVAVAV-------------ILMVLGLLTIGSIFFTWRLYNER-------
BTF1    PESFMPSVSPCAVALP-------------IIVVILMIPIAVCIYWINKLQKEKKILSGEK
BTF2    PESFMPSASPWMVALAVILTASPWMVSMTVILAVFIIFMAVSICCIKKLQREKKILSGEK
BTF5    ADPFFRSAQRWIAAALAR-------------TLPVLLLLLGGAGYFLWQQQEEKKTQFRKK
BTF3    ADPFFRSAQPWIAALAG-------------TLPISLLLLAGASYFLWRQQKEKIALSRET
BTF4    ADPFFRSAQPWIAALAG-------------TLPILLLLLAGASYFLWRQQKEITALSSEI
                       *.                 .  .         .        *


BT      PRER----------RNEFS-------SKERLLEELKWKKATLHA---------------
BTF1    EFERETREIALKELEKERVQKEEELQVKEKLQEELRWRRTFLHA---------------
BTF2    KVEQE---------EKE---------IAQQLQEELRWRRTFLHA---------------
BTF5    KREQELREMAWSTMKQEQS-------TRVKLLEELRWRSIQYASRGERHSAYNEWKKALF
BTF3    ERERENKEMGYAATEQEIS-------LREKLQEELKWRKIQYMARGEKSLAYHEWKMALF
BTF4    ESEQEMKEMGYAATEREIS-------LRESLQEELKRKKSST----------------
              *.              *              * ***. .


BT      --VDVTLDPDTAHPHLFLYEDSKSVRLEDSRQK---LPEKTERFDSWPCVLGRETFTSGR
BTF1    --VDVVLDPDTAHPDLFLSEDRRSVRRCPFRHLGESVPDNPERFDSQPCVLGRESFASGK
BTF2    --ADVVLDPDTAHPELFLSEDRRSVRRGPYRQR---VPDNPERFDSQPCVLGWESFASGK
BTF5    KPADVILDPKTANPILLVSEDQRSVQRAKEPQD---LPDNPERFNWHYCVLGCESFISGR
BTF3    KPADVILDPDTANAILLVSEDQRSVQRAEEPRD---LPDNPERFEWRYCVLGCENFTSGR
BTF4    ------------------------------------------------------------


BT      HYWEVEVGDRTDWAIGVCRENVMKK-GFDPMTPENGFWAVELY-GNGYWALTPLRTPLPL
BTF1    HYWEVEVENVIEWTVGVCRDSVERK-GEVLLIPQNGFWTLEMH-KGQYRAVSSPDRILPL
BTF2    HYWEVEVENVMVWTVGVCRHSVERK-GEVLLIPQNGFWTLEMF-GNQYRALSSPERILPL
BTF5    HYWEVEVGDRKEWHIGVCSKNVQRK-GWVKMTPENGFWTMGLTDGNKYRTLTEPRTNLKL
BTF3    HYWEVEVGDRKEWHIGVCSKNVERKKGWVKMTPENGYWTMGLTDGNKYRALTEPRTNLKL
BTF4    ------------------------------------------------------------
```

Figure 3 (Page 1 of 2)

```
BT      AGPPRRVGIFLDYESGDISFYNMNDGSDIYTFSNVTFSGPLRPFFCLWSSGKKPLTICPI
BTF1    KESLCRVGVFLDYEAGDVSFYNMRDRSHIYTCPRSAFSVPVRPFFRLGC-EDSPIFICPA
BTF2    KESLCRVGVFLDYEAGDVSFYNMRDRSHIYTCPRSAFTVPVRPFFRLGS-DDSPIFICPA
BTF5    PKPPKKVGVFLDYTGDISFYNAVDGSHIHTFLDVSFSEALYPVFRILTLEPTALSICPA
BTF3    PEPPRKVGIFLDYTGEISFYNATDGSHIYTFPHASFSEPLYPVFRILTLEPTALTICPI
BTF4    ------------------------------------------------------------


BT      ADGPERVTVIANAQDLSKEIPLSPMGEESAPRDADTLHSKLIPTQPSQGAP---------
BTF1    LTGANGVTVP---------------EEGLTLHRVGTHQSL--------------------
BTF2    LTGASGVMVP---------------EEGLKLHRVGTHQSL--------------------
BTF5    ------------------------------------------------------------
BTF3    PKEVESSPDPDLVPDHSLETPLTPGLANESGEPQAEVTSLLLPAHPGAEVSPSATTNQNH
BTF4    ------------------------------------------------------------


BT      ----------
BTF1    ----------
BTF2    ----------
BTF5    ----------
BTF3    KLQARTEALY
BTF4    ----------
```

Figure 3 (Page 2 of 2)

FIG. 4A.

FIG. 4B.
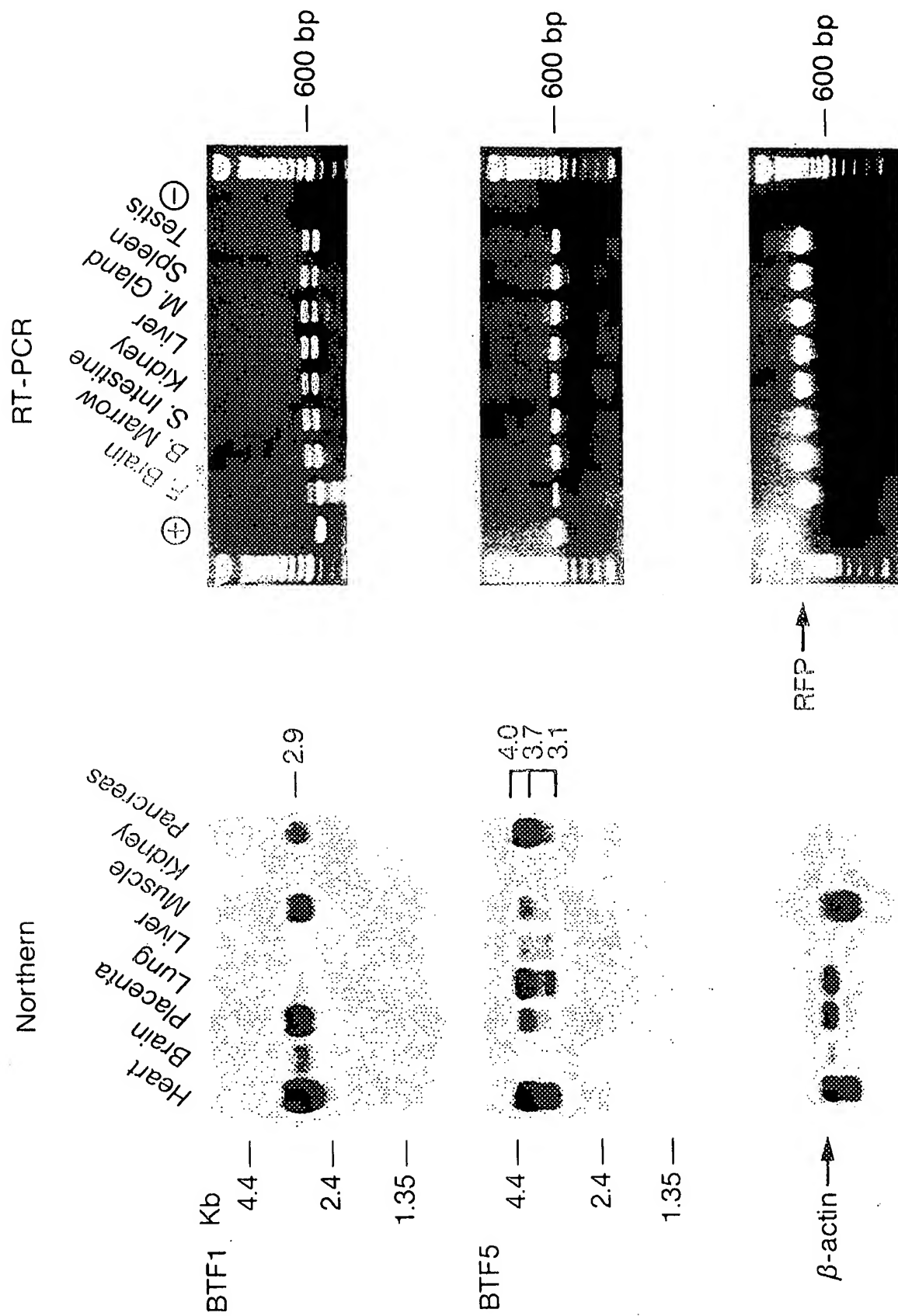
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                                        ┌─────────CYSTEINE-RICH DOMAIN─────────
52 kD Ro   MASAARLTMMW│EEVTCPICLDPFVEPVSIECGHSFCQECISQVGKGGGS-----VCPVCRQRFLLKNLRPNRQLAMMVN
RoRet      MASTTSTKKKM│MEEATCSICLSLMTNPVSINCGHSYCHLCITDFFKNPSQKLRQETFCCPQCRAPFHMDSLRPNKQLGSLIE
           ***  *   **│*  ** **** *  **  **** *****  *  **    *

52 kD Ro   NLKKISQEAREGTQGERCAVHGERLHLFCEKDGKALCWVCAQSKKHRD┌HAMVPLEEAAQEYQEKLQVALGELRRKQELAEKL
RoRet      ALKKTDQEM------SCEEHGEQFHLFCEDEGQLICWRCERAPQHKGH│TTALVEDVCQGYKEKLQKAVTKLKQLEDRCTEQ
           ***       *   * ***  ***** *  * ** *      *   * │   * *  *  ***** *  * **  **

52 kD Ro   EVEIAIKRADWKKTVETQKSRIHAEFVQQKNFLVEEEQRQLQELEKDEREQLRILGEKEAKLAQQSQALQELISELDRRCHS
RoRet      KLSTAMRITKWKEKVQIQRQKIRSDFKNLQCFLHEEEKSYLWRLEKEEQQTLSRLRDYEAGLGLKSNELKSHILELEKKCQG
               * *  **    * *  * * * *  * **  *          *  * *               *  *     * *

52 kD Ro   SALELLQEVIIVLERSESWNLKDLDITSPELRSVCHVP-----GLKKMLRTCAVHIT┌LDPDTANPWLILSEDRRQVRLGDTQQ
RoRet      SAQKLLQNVNDTLSRSWAVKLETSEAVSLELHTMCNVSKLYFDVKKMLRSHQVSVT│LDPDTAHHELILSEDRRQVTRGYTQE
           ** ***   * * *   * *  * *   * *   *         ***** *     │*****   * *********  * **

                                        ─────────B30-2 DOMAIN─────────
52 kD Ro   SIPGNEERFDSYPMVLGAQHFHSGKHYWEVDVTGKEAWDLGVCRDSVRRKGHFLLSSKSGFWTIWLWNKQKYEAGTYPQTPL
RoRet      NQDTSSRRFTAFPCVLGCEGFTSGRRYFEVDVGEGTGWDLGVCMENVQRGTGMKQEPQSGFWTLRLCKKKGYVALTSPPTSL
               * ** *  ***  * *** ** **** *    *****     *  *     ****  *     *        *  **

52 kD Ro   ┌HLQVPPCQVGIFLDYEAGMVSFYNITDHGSLIYSFSECAFTGPLRP┌FFSPGFNDGGKNTAPLTLCPLNIGSQGSTDY
RoRet      │HLHEQPLLVGIFLDYEAGVVSFYNG-NTGCHIFTPKASFSDTLRP│YFQVYQYS-----PLFLPPP--G----D---
           │**    *  ********* ****   *   * *     *  *  ││  *            *  *        *
```

FIG. 5A.

```
NPT1   MQMDNRLPPKKVPGFCSFRYGLSFLVHCCNVIITAQRACLNLTMVVMVNSTDPHGLPNTSTKKLLDNIKN----------
NPT3   --MDGKPATRKGPDFCSLRYGLALIMHFSNFTMITQRVSLSIAIIAMVNTTQQQGLSNASTEGPVADAFNSSISIKEFDTK
NPT4   MQVDETLIPRKGPSLCSARYGIALVLHFCNFTTIAQNVIMNITMVAMVNSTSPQSQLNDSSE-----------

NPT1   -PMYNWSPDIQFIILSSTSYGVIIIQVPVGYFSGIYSTKKMIGFALCLSSVLSLLIPPAAGIGVAWVVVCRAVQGAAQGIVA
NPT3   ASVYQWSPETQGIIFSSINYGIILTLIPSGYLAGIFGAKKMLGAGLLISSLLTLFTPLAADFGVILVIMVRTVQGMAQGMAW
NPT4   ----------------VLPVDSFGGLSKAPKSLP--------------------AKSSIL

NPT1   TAQFEIYVKWAPPLERGRLTSMSTSGFLLGPFIVLLVTGVICESLGWPMVFYIFGACGCAVCLLWFVLFYDDPKDHPCISIS
NPT3   TGQFTIWAKWAPPLERSKLTTIAGSGSAFGSFIILCVGGLISQALSWPFIFYIFGSTGCVCCLLWFTVIYDDPMHHPCISVR
NPT4   GGQFAIWEKWGPPQERSRLCSIALSGMLLGCFTAILIGGFISETLGWPFVFVFIFGGVGCVCCLLWFVVIYDDPFSYPWISTS

NPT1   EKEYITSSLVQQVSSSRQSLPIKAILKSLPVWAISIGSFTFFWSHNIMTLYTPMFINSMLHVNIKENGFLSSLPYLFAWICG
NPT3   EKEHILSSLAQQPSSPGRAVPIKAMVTCLPLWAIFLGFFSHFWLCTIILTYLPTYISTLLHVNIRDSGVLSSLPFIAAASCT
NPT4   EKEYIISSLKQQVGSSKQPLPIKAMLRSLPIWSICLGCFSHQWLVSTMVVYIPTYISSVYHVNIRDNGLLSALPFIVAWVIG

NPT1   NLAGQLSDFFLTRNILSVIAVRKLFTAAGFLLPAIFGVCLPYLSSTFYSIVIFLILAGATGSFCLGGVFINGLDIAPRYFGF
NPT3   ILGGQLADFLLSRNLLRLITVRKLFSSLDMQVSSWE----------------SQGDLGSSQES-SLPLPLDSSS-----
NPT4   MVGGYLADFLLTK-KFRLITVRKIATILGSLPSSALIVSLPYLNSGYITATALLTLSCGLSTLCQSGIYINVLDIAPRYSSF

NPT1   IKACSTLTGMIGGLIASTLTGLILKQDPESAWFKTFILMAAINVTGLIFYLIVATAEIQDWAKEKQHTRL
NPT3   ---VRILSLVGGMSFSCLL-----QSTCLAWSFTSRLDKQNFKTGPKRGPLPASEDIKLQT------
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FIG. 5B.
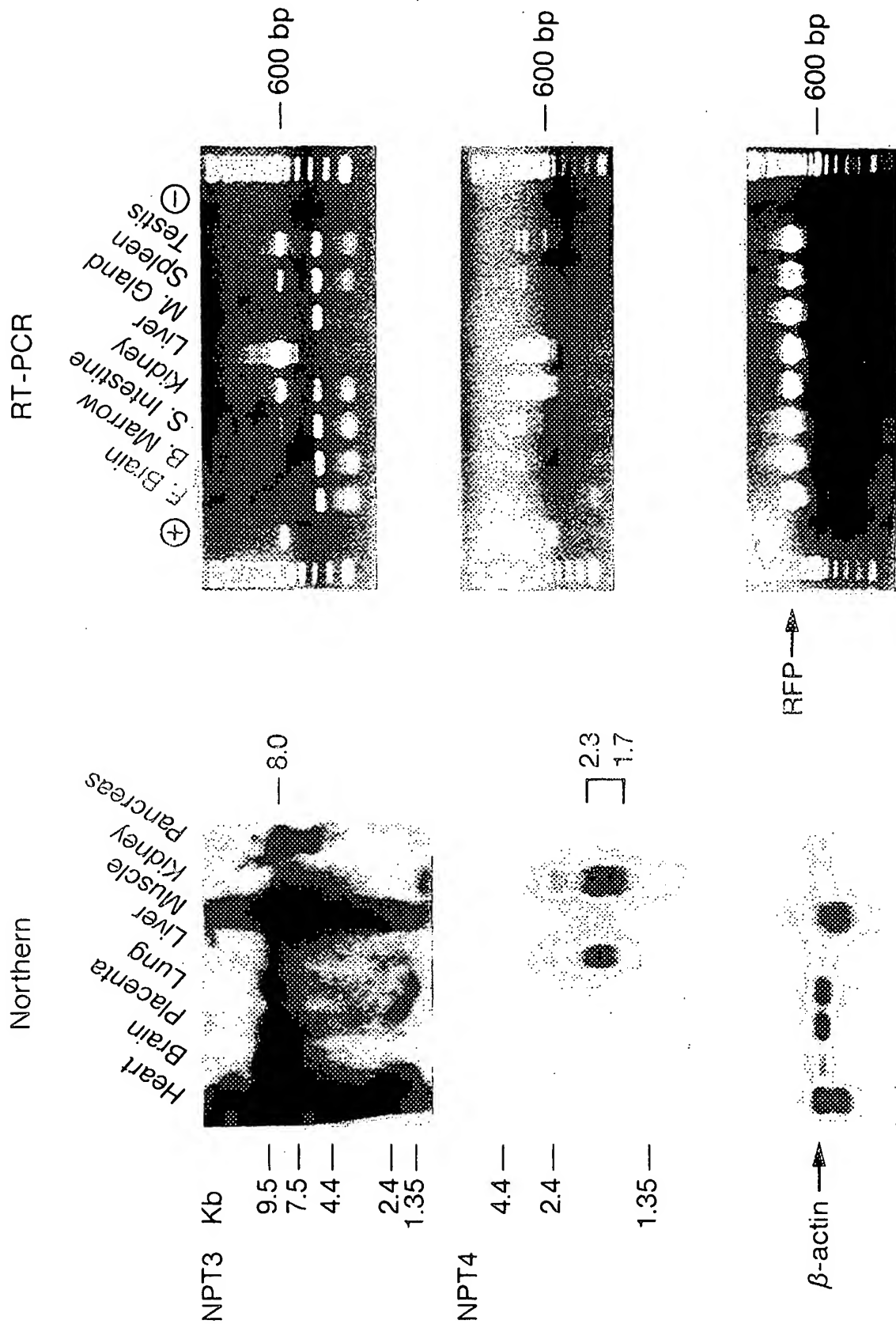
RT-PCR

FIG. 6B.

FIG. 6A.

Northern

FIG. 6D.

RT-PCR

FIG. 6C.

Northern

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Figure 7 (1 of 6)

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Figure 7 (2 of 6)

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Figure 7 (5 of 6)

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Figure 7 (6 of 6)

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3541  TGGGATGGAA  TAATTTTTTT  TTTTTTTTTG  AGACGGAGTC  TAGCTCTGTC  GCCCAGGCTG
3601  GAGTGCAGTG  GCACCATCTT  GGCTCACTGC  AAGCTCTGCC  TCCCGGGTTT  ATGCCATTCT
3661  CATGTCTCAG  CCTCCAGAGT  AGCTGGGACT  ACAGGCGCCC  GCCACCACGC  CTGGCTAATT
3721  TTTTTTTTAT  TTTAGTAGAG  ATGGGGTTTC  ACCGTGTTAG  CCAGAATGGT  CTCGATCTCT
3781  TGACCTTCTG  ATCCGCCTGC  CTTGGCTTCC  CAAAGTGCTG  GGATTACACG  TGTGAGCCAC
3841  CATGCCCGGC  TGGGATGGAA  TAAATTTATC  TTGTATGGGA  GAAGGACATA  CATTTTGGCA
3901  GGTCAAGGAC  AGAATGTTAT  GGACTAAACT  GTGTCCCCCA  AAATTCATTT  ATTAAAACCC
3961  TAAACCCCAG  TGTGACTGCA  TTTGGACATA  GAGCCTTTAG  GGGGTACATA  AAACTAAAGA
4021  TCACAGGATA  GGGCCCTAAT  CCCATTGGGG  CTGGTGTCCT  TACAGAAGAT  GAGACACTTA
4081  GAGCTCTCTC  TCCACGCAGG  CACCAAGGAA  ACACCATACA  AACACACAGT  GAGATGGCAG
4141  CCATCTGTTA  GCCAGGAACA  GATTCTCACC  ATAAACTATG  TTGGCACCTT  GATCTTAAAC
4201  TTCCAGGCTC  CAAAACTGTG  AGAAAATGAA  TTTCTGTTCC  AAGCCTCTTA  GATATGGAAA
4261  AAAAGATTCT  GTTGTTTAAG  CCATCCAGTC  TCTGGTATTT  TGTTATGGCA  GCCTGAGTAG
4321  GCTAAGACAA  TGAAGGATGT  GGTAAAACTT  TACGTCCCAA  CCACATACCA  AAGAGGCTGG
4381  AATTTAGCAT  GCTTTCTTCT  TTCAACTGTA  GGCAATGTGC  ACAAGTTCTA  AATCCTAAGA
4441  CATGTTGGCT  CCTTTACTCT  GCCCAAACTA  CAACTCAAAC  AAACAACTGT  AATATAATAA
4501  CATCCAATGA  AGTTCTGACA  TTTCTTCAAC  ATGAGTACAG  TAATTCAATG  CCAGAGAATT
4561  CATTTTATTT  TGAAATCTAC  ATGCCATATT  CCAATTTCTG  TTGAAGATGC  AATGGTTATA
4621  TTTATTCTTT  TTAATATAGA  TTTATCAGAC  TGGGCGCGGT  GGCTCATACC  TGTAATCCTA
4681  GCATTTGAGA  GGCTGAGGTG  GGCATATCAC  CTGAGGTCAG  GAGTTTGAGA  CCAGGCTGGC
4741  CAACATGGTG  AAACCCTGTC  TCTACTATAA  ATATAAAAAT  TAGCTGGGTG  TGGTGGTGCA
4801  TGCCTGTAGT  CCCAGTTACT  AGGGAGGCTG  AGGTAGAATT  GCTTGAACCT  GGGAGCAGGA
4861  GGTTGCAATG  AGTGGAAATC  GCACCAGTAC  ACTCCAGCCT  GGATGACAGA  GCAAAATAAT
4921  AAATAAATAC  ATAAAATAGA  TTTATCAGTT  TATCAATAAT  ATAGTTTTCT  TTTCTAGGTG
4981  TAAATATAGG  TAATGACTGT  CCTTTAGTAC  ATTTTCTCAT  GATGCTCCTC  TTACTTGGTT
5041  TGGTACAATA  TTAAGTATTG  AAATAAAATA  GAGAATCCTG  TCGCTACACA  TGAGCACTTA
5101  TTCCATTTGC  TCATCTCCAA  TATGCACGGG  AAATTCTCAA  ATTGCTAATA  ATCTTGTAAC
5161  ACACATGCAT  TATATTCAAC  AGGAATATAT  AAATTTATAA  TTATAATTTA  GGATCAACAG
5221  ATGACAAACC  TTTAGAAGGT  TTGTATTTAA  CCTTAAAATA  TAATTTTTTA  AAAATTGGTT
5281  ATAAAATTTC  TAATACTTTC  TTTTTTGTGA  CCTCAAGGGG  AAAATATAAT  TCTTATAAAA
5341  GTTCAAATGA  TTTACAGAAT  ACAAAAGTG   AATAGAGATG  ATGAATGAAT  TAAAGGAAAG
5401  GATATTGCTA  CATAGATTTG  GAAATTTAAA  AAGGGAAATT  ACGATTGTTG  ATTTTGTGTT
5461  AAACTGATCT  GCTTTGTTCA  AGATACCTTA  TGTACCAAAA  AATGATTTTA  TCTCAGCCTC
5521  ATATCTCAGT  AAATTCCTGA  GACAAACTTT  AGTCCCTGGT  GCCCAGGTGC  CTTTGGTAAT
5581  TGGGAGACCT  CTAGGTTTAG  CATCCTCATC  CACTCGCCCC  AATTTAAATA  GTCCTCCCCA
5641  GGGCCATTCA  GGCAAGGGAG  ATGAAAACTT  GCTCAAGAGT  TGGAATCCAA  CTGAAGCTAC
5701  CGAAATTCAT  TGCTCAATAG  ATAATTTTCC  CTGGAAGTAA  CTAGGGCTTT  TGAATATAAT
5761  AGTGGGCATT  TCAAAGTAGA  AGGTAAAGTA  TTTTGGAGAT  GAGGAGACAG  GACAGAGCTA
5821  CGAGGAATGT  CCTTTGCTTA  GGGACTAGGC  TCTTAGCAGT  ACCTCTTAGG  TAAGAACTGG
5881  TTAACTGGCA  CCTTCTGTGT  TTCTCTGAAG  CTCCCTTTGC  TTAGGGACTA  GGCTCTTAGC
5941  AGTACCTCTT  AGGTAAGAAC  TGGTTAACTG  ACACCTTCTA  TGTGTCTGAA  GCTCCCAGAA
6001  CAAACTGCCA  GTGAAATTTG  GATTTTTGGA  ATATAGTTTC  TTTTTTCTTG  TTACTTTTTG
6061  TTTTGTTGTT  TTTTTTTGAG  AGTCTCACTC  TCACTGCAAC  CTCCCCCTCC  TATATTCAAG
6121  TGATTCTCTT  GCCTCAGCCT  CCCGAGTAGC  TGGGACTACA  GGCGTGCACT  AGCATGCCCA
6181  GCTAATTTTT  GTATTTTTA   GTAGAGATGG  GGTTGGTTTT  TTTTTGAGAC  GGAGTTTCAC
6241  TTTGTCGCCC  AGGCTGGAGT  GCAGTGGCAC  GATCTTGGCT  CACTACAACC  TCCACCTCCC
6301  GGGGTTCAAG  TGATTCTTCT  GCCTCAGTCT  CCTGAGTAGC  TGGGACTACA  GGCGCCACA
6361  GGTGAACACC  GCCACACCTG  ACTAATTTGT  GTAGTTTTAT  TAGAGATGGG  GTTTCGCCAT
6421  GTTGGCCAGG  CTGGTCTCAA  ACTCCTGACC  TCAGGTGATC  TACCCACCTC  AGCCTCCCCA
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6481  AGTGCTGGGA  TTACAGATGT  GAGACACCAG  ATCAGCCTCA  GAAGACATTT  TCTATTGGAA
6541  AGAGAAAACA  CTATTAGCAA  CCTATTAGTC  TAATATTTAA  TACTTAATGT  CTTCCTTAGT
6601  AATAAACCAA  CTCTCTACAA  CAAAGTGCTT  CCTGGCTGCC  TAAGTCATTG  ATTCATTCAG
6661  TTCAACATTT  TCTCAATGCC  CAACAGCCAA  GTGTCTCTTG  TATGCCAAGT  TCTATGCTGA
6721  TTATCAGTAT  TTGAATAAGA  GGGGGTCTAC  ATCTTAAGTA  CTGCTTAAGA  TGAAAGCCTC
6781  TAGGTTAACA  AACTTAACAC  AATGTATCAT  TCACTACTAA  ATAGACCGAA  TACAAAATCT
6841  TGTTATTGGA  GCCCAGAGAG  AAGAATTGAA  ATTCAAGTTT  TCTCTCTCTC  CTTTTCTCAC
6901  TCACCACAAT  AAGTCAGTTG  CACCAAGTCT  TGTAGCTCTT  TACTGAGCCA  TGTTTTCACG
6961  TGTCCCTTTG  TTTTATTTGC  CACACCCTAA  ATAAAAATTG  TACTGGCTTT  TTTTCCCTGG
7021  GTTTACAGTA  TTAATACATT  GTCAAGATTT  ACCTCTTCGT  GTAGATTCCC  TGGGGAAAAT
7081  TACCTTTCCT  CCTTCCCTTA  AATTCTTCAG  AGGTTAGAAA  GCCATTAGTA  ACATTCTGGT
7141  ATGTGGACAA  AGTTACCCA   TTATGTATGG  ATGTTTTACT  CTTTCTATTT  TTCTGACAAT
7201  AATCTCTTAA  GGAGGTGTGG  TTATAGAATA  GTCAGCTGTT  ATAAGTACTG  TTTTCCTGGC
7261  CTTACAACTT  AAGTTCTTTA  AGCTGTTTCT  TAGTTTGCTC  ATCTCAAAAT  TCGGAATAAG
7321  GATAAAACCT  ATCTCTTAGA  TTGTTGGATT  AAATGAATTA  ACATACTGGA  AGCTCATGAA
7381  ATGTGCCTGG  CACACAGTAG  TGCCTAATAA  ACCATCTCTC  TTATTCAGCC  TGTTTTCTGA
7441  TTTCAGAATC  TACACTTGCT  GAGCCAGGTT  CTTTTCATTT  CAAGGTGAGC  AAAAGCATAC
7501  AAGGAAGAGA  TGGAGGTAGG  AAGAGATTAA  GCCCTAGGCC  AAGGTCACAC  ACCGATTGGG
7561  AGCTGGAATC  AAAGGCAATT  TGGTCAGTGA  ATAAAAAGGA  TTCCAAGGCC  CATAAGGCAA
7621  TTCTAACCTT  AGGATCGAAA  TTCTCGGACA  TACAGGAAAT  GCTGGGGGGG  GAAAATCCGG
7681  TCTTCTCAGC  CCAAGAGCCA  TGTGAAACCA  GACCTTCAAA  TCTGATGATT  CTCAGCCCAG
7741  CTGCCCATTA  GAATCGTTGT  AATTTAAAAA  TACCCTCGGA  AAATTCTAAT  ATGTGGCTAT
7801  CAAAGGTGAT  CATTTGCTTT  TATGCCACTT  TGTTTTCACC  CAAATGGGAC  ATCCAACCCT
7861  TTTCCTTTGA  GAGTAGTTGT  AGGGAAAGGA  GGGGGTGGAG  GGAGGGAAGA  GCGGAAAAGG
7921  CTGGATCCGC  CCTGAGCCGG  TGTCAGTATC  TGGGAAGTGG  GAGGCGCGTC  AGCAGTAAAC
7981  AGCTTCTGCT  AGGATTATTA  TCTCCTGCCA  CACACTCGGA  TTTGAAGGCT  CCAAACGAAA
8041  CAATGCAAAA  CGCTTCAGTG  GAGTTCCAGA  AGCGTTAGAC  TAAACGACTG  GGTCTGTTTG
8101  GCCAGTCTGA  GCAGCTGGGC  GCAGATGCAT  AGGCAAGACT  TAGCCCGCCT  AGACTTTTCT
8161  GCCCACTTAA  TTCCGATCAA  AGCAGAAACC  GGCCGGGCGC  GGTGGCTCAC  GCCTGTAATC
8221  CCAGCACTTT  GGTAGGCAGA  GGCTGGCGGA  TCACCTGAGG  TCAGGAGTTC  GAGACCAGCC
8281  CGGCTAACCT  GGTGAAACTC  CGTTTCTACT  GGTGGCGGGC  GCTTGTAATC  CCATCTACTA
8341  GGGAGGCTGA  GGCCGGAGAG  TCGTCTGAAC  CCGGGAGGCG  GAGTTTGTAT  GCAGTGAGCC
8401  GAGATCGCGC  CACTGCATTC  CAGCTTGGGC  AACAGGAGCA  AAACTCCGTT  TCAAAAAAGC
8461  AAGCAAACAA  ACAAAAAAAT  GCAGAAACCG  AGATCCGGAA  GAAAACCTCG  GCGAGATTCA
8521  CAGAATCCAG  GAAAATAGGT  CTCTAGAAAT  TTGTCCATGG  TCCCAGATCT  CCATTTCTTG
8581  TGGGTGGGGC  AGCTGTTACC  AGATCCCTAG  AAGCAAAGGT  TTTTTTGGGG  GACCGTGTCT
8641  CACTGTTGCC  CAGGCTGGAG  GGCAGTGGCA  CGATCTCGGC  TTACTACAAC  CTCCGCCTCC
8701  CAGGCTCAAG  CGACTCTCCT  GCGTCAGCTT  CAAGAGTAGC  TGGGATTACA  AGGTATGTGC
8761  CACCACGCCC  AACTTATTTT  TTTATTTATT  ATTTTTATTT  AGTAGAGAGG  TGTTTCACCA
8821  TGTTGGCCAG  GTTAGTGTCG  AAGTCGTGAC  CTCAGGTGAT  CAGCCCCCTC  GGCCTCCCAA
8881  AGTGGTAGGA  TTAGAGGGGG  GAGCAGAAAG  CAAAGGTTTT  TGAGTGGCCA  CAGGCCCCAC
8941  TCTATTTCCT  TTTCTGCCTG  TAATGGCAAC  CTAGACGCTT  GAGCTTCTTA  AAATACAAGA
9001  GTAAGTTGCA  TGTCAGGCAC  CGTTCTACAT  TAGGGACATT  AGTCTGTTTT  ACAGACACCT
9061  TTCAACTCCC  TGGTTAACTT  TTAGGTAATA  TACTCTGCAC  TTTAGCAGGA  ATGGGACCTA
9121  TAACTCTCAC  AGAATTAGGA  AAGTGAGGCT  GCCTACAGCC  TAAATTGAGA  AAAAAATAGA
9181  CGGGGGACTA  GTCGGAGGAC  CAAACAAGGT  TACCAACACG  TTAGAGTTTT  GCCTTCAATT
9241  TACATTTTTA  AAGTAATCAC  AACGAAGTGT  TTAGATCACG  AGGCATCCCT  GCATGTAAAC
9301  TGTTAGGCAC  TAACTATGGT  CGATCTTACA  AAGCATTAAC  TAGAATATTT  CTTTAGAGTA
9361  TGATAGTACG  TAACTGACCT  ACTATTACAT  ACAAACAGAC  CAACCTTTAG  TAACAGCGCT
9421  CCCCAAAAAC  CGAAAAGCAG  TAATACGCTT  TGCTCAAGGT  TGGCATAAAA  TTAACTTACC
9481  TTAGTGCCTT  TTTTCCTTCT  ACCTACAAGC  AGTGAGGTTA  GCTCTTCCTT  TGAAACGGTA
9541  GGGGGGCTCT  GAAAAGAGCC  TTTGGGTTTG  ATAGCGTTTC  CGGGAGCTCA  GATACCTGTC
9601  AAATCACTTG  CCCTTGGCCT  TGTGGTGACT  CTCGGTCTTC  TTAGGCAGAA  GCACGGCCTG
9661  GATGTTAGGA  AGGACGCCGC  CCTGAGCAAT  GGTCACCCGG  CCTAGCAGTT  TGTTGAGCTC
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 9721   CTCGTCGTTG CGGATGGCCA GCTGCAAGTG GCGCGGGATG ATGCGAGTCT TCTTGTTGTC
 9781   GCGAGCCGCG TTGCCGGCCA GCTCCAGGAT CTCGGCGGTC AGGTACTCTA ACACCGCCGC
 9841   CAGGTACACC GGCGCGCCTG CCCCAACCCG CTCTGCGTAG TTGCCTTTAC GGAGCAGGCG
 9901   GTGCACTCGG CCCACCGGGA ACTGGAGACC AGCGCGAGAA GAGCGGGATT TCGCTTTGGC
 9961   GCGAGCTTTG CCTCCTTGCT TACCACGTCC AGACATTGCA ATCAGACAAA AATCACCAAA
10021   ACCAGCGGCC TAAGCTCACG AGAAACAAA  CAAAATCAAG AAATATGTAA AACATGGCCG
10081   CTTTTATAGG TAGTTCCTGG GGAGTAAATC CGACTTTTTG ATTGGTCGGT AGCAAATGCT
10141   AGTCAGATAG CCAATAGAAA AGCTGTACTT TCATACCTCA TTTGCATAGC TCTGCCCACG
10201   GATGACAACT GTGCAGTTTG TCTTCCAATT AACTAAGAGG TACTCTCCAT CCCTCATTAG
10261   CATAAAAGCC CTATAAGTAG CAGAAATCCG CTCTTTACTT TCGACACATT TCTGGTGTTT
10321   TAAGATGCCT GAGCCAGCCA AGTCTGCTCC CGCCCCGAAG AAGGGCTCCA AGAAGGCAGT
10381   GACCAAAGCG CAGAAGAAAG ATGGCAAGAA GCGCAAGCGC AGCCGCAAGG AGAGTTACTC
10441   TGTGTACGTG TACAAGGTGC TGAAACAGGT CCATCCCGAC ACTGGCATCT CTTCCAAGGC
10501   CATGGGCATC ATGAATTCTT TCGTTAACGA CATATTTGAG CGCATCGCGG GCGAGGCTTC
10561   CCGCCTGGCG CATTACAACA AGCGCTCGAC CATCACCTCC AGGGAGATCC AGACGGCCGT
10621   GCGCCTGCTG CTTCCCGGAG AGCTGGCCAA GCACGCCGTG TCGGAGGGCA CCAAGGCCGT
10681   CACCAAGTAC ACCAGCTCCA AGTAAACATT CCAAGTAAGC GTCTTAACAC CTAACCCCAA
10741   AGGCTCTTTT AAGAGCCACC CAGATACCCA CTAAAAGAGC TGTGGCCAGA CGCCAAATTT
10801   TATTTGGCGG CGGAGGGGTA TTAGAATATA GGAACTGGAG AGGGGTGGGG ACAAGTGTTG
10861   CAGCTTAGAG AGGGACAAAG GGTCCTGAAC CCGAAAGAAG CCAGCCATTA AAAATGGCTT
10921   TGGGGTCAAT TCGTTGTGCT TAAATTTAAA ATGGAGACAA GCGGCCATTT TGCTAACTCG
10981   GCGTTCCCGG AAGAAACCGC AGGCTCGCTT AGGTTTCAGA CCCAGCTGTC TGTCCCTGTC
11041   TACGTCGCCA GGATCAACGG TTGCCGTAAT GTCATAATTT CGCCACCAGC TTCTAGCCAA
11101   TAGGCTGTCC TGTCATTTTA AATATTAACC AATCGAGGGA AAGCTGTTTT GAGACTCTGA
11161   TTTACATAGC GGACCGGAGT GGGAACCTGG GCAGTAACTG CCTAAGGAAG GACTCCCCCT
11221   CTGTTTTCGT GGCGCACACC TTCGTAGTAT ACTGAAGGGT GTGTCTCCTG GGTTTCCAAC
11281   TGCCCCGGTA ATAGTCTTTT AACCTAATAT GCGTCAGTTT TGATAACAAC ACTAAGGCAG
11341   TACAGAACTA AAGATGTAAG CACTGCGCCA GATGTTGCTT CATACATCTT ATTCTATTCA
11401   ACTGGTTTAT TCAAGATTCA AATCAAATCA AATTTTGCTT GAATCCCAGT GCTCAGTCAG
11461   CCATAAATGG TGTGTTGCCT GATTGAAACT TAAAATCTCC GTAGGGGGCT TGTAACATGC
11521   AGACAAGTTT GAAAGTTGCT TTAGGAGAAG CCAACTCTTA ACTGCTGGGT AAATTGACAA
11581   GCCTTCGAAC ACTGAACTGA AGGCCAGTAA GGACTAGGCG CTGGGTGGGG GAGAATGAAG
11641   AGGAGACGTC ATTAAACTTA GCACATACAC TGTATCTCCT AGAGGACTCT CCCTTCCTAG
11701   ACAACTGCAG GCCGCTTTGT GGCCTGGGAA ATTCCACATT CCCTTAAGTA TTTTACTCAT
11761   GGTCTTTTCC AGGTAAAGAT TTTAAGATGA AGGGTTAGAC GTAGTCTACC TATCTTTTTA
11821   TTCAAGTCTA GAACACGTTT TTAGCACCTA GAAGTTTGCT TTCTCCATTA AAAACCGGGA
11881   ATATACAATA AATAAAATTA GTGTTAAAGC AGATTTTTAC AAACTTAAAT ACCATGTAAT
11941   TTAGGTTACA GTTATTTAAC ATAAGGACTG TGTGATCTTA AATCTGCAAT TTCTTTCACA
12001   CCTGGGAAAT AAACTAAGGC CTGTCTTTGG TGCCAGACAA GGCCTTATAC TTGAACACTG
12061   CTGTGCAATC ACAGGCTGCC TTGCCTAGAT AACTTATCTG AGAAATTCTG ATGAGAAATG
12121   AAATTTCCAG AGTCCCTCAC AAGTAAATTT TTTTTTCTTT TTTTTTTTTT TTTTGAGAC
12181   GAAGTTTCTC TCTTGTTTCC CAGGCTGGAG TGCAATGGCG CGATCTTGGC TCACAGCAAC
12241   CTCCGCCTCC CGGGTTCAAG CCATTCTCCT GCCTCAGCCT CCGGAGTAGC TGGGATTACA
12301   GGCATGCGCC ACGACACCCT GGCTAATTTT GTATTTTTAG TAGAGACGAG GTTTCTCCAT
12361   GTCGGTCAGG CTGGTCTCGA ACTCCGGACA TCAGGTGATC TGCCCGCCTT GGCCTCCCAA
12421   AGTCCTGGAT TACAGGCTTG AGCCACCGCG CCGGGCCTAA ATGGTTTTTT TTTTTTCTAT
12481   GCCTCTAATG GACCTGGTCA CTTATTCCCA TTCAGACTGA CCGCTCTCCT ACCTGCCAAC
12541   TAACTAATCA GTGTAACCAA AATCTGCAAA CAAAATTCAG TATTCTTTCC CCGCCTTTTC
12601   CCCTTTCTCT TACATAGATT ATGTTTTTGC CTGTGTTAGA TGAAATAATT CTATTGCTTG
12661   TTCTCTCTTC TGTACAAGTA CCCAGTAAGC AAATTATTAA CTTCTTGGTC ATTTATTTCT
12721   GAATTTTCCA CCAAGACAGT GTTTATGTGA GTCATACAAT AAGAACCAAC AGAAATGTGT
12781   GTCTTGGAAA CAGGTTGTCT ATCCCTGGAC CCTTTGAGTT TTCTGTTCAC TTTCCTTTGG
12841   CTTTTGCATG CTAAAAGTTT ATCGTCCGCG TTTGTTTGTT TTGGTTATTC TAATTGGACT
12901   TGGCTGATTG GTTGCATATT GGTGGCAGTA GTAGAATTTG AATTCTGGTT TTCTGGTCAC
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12961  ATCATTAAGT  GATTAGTCAG  TGGAGAGGAC  AGGAAATCTG  GTTTATTTAT  TAACCTTTTT
13021  TTGGGGTGTT  TTTGTTTGAA  GATGTTGATA  TTCTCTGTGA  GGACACAGGG  TTAGAGTTGG
13081  TGTTTTTCTT  TCTGACTTTA  CATGGGATTT  GATGTTTTGT  GCTTGTATGC  CTCTTTCCAC
13141  CTTCCAAAAC  TTGTCTTTTT  TGAGTCCAAA  TAGTTGTCGA  TATCTGCAAA  ACCAGTATTC
13201  CTGTGTTAAG  ATGATATGAA  TATAAAATGG  CTGCCCTGTT  ATAACTTTTG  ACTTTAAGAA
13261  AGTGTTAGGA  CTAACAGGAG  ACAAAAAGGA  AATCAAGGAA  ACCGAATGTC  TGGTCTCAAT
13321  AACTGCTATG  GCAGAGGCTC  TACAGCTTAT  TATTAATTTT  AGTAATTTCA  CATTATTGCC
13381  CCTTCACGTT  CTTTAAGTAA  GGTTAGAGGA  CAGAAGAAAC  ATAATGTTGT  TACAAATTGG
13441  ACTATTGAGT  CAGGGAAAAA  AAAGAGTGCT  TTCAATATCT  GAATAAAACA  AAGATTTAAT
13501  ATTTTCTAAA  CCTTAACGAG  TTTATTGTAA  GGGATGTGAT  GCTGGAAACT  AGGAAACTAG
13561  AATTTTCTTC  TAAACTGAGA  ATCAGAATTA  TTCATATTCT  CAGCAGTGGT  GCCACCTGAG
13621  GGACTTCTGA  TCTTAATTAC  ATACTTTTAT  TTCTTTAACT  GATCAACATG  CTAAATAGAT
13681  AACCTATGGC  TCTGTTTTTA  CCCACTTTAA  ATTCTGTTCT  ATTAGCACGG  TTAGCTTTCC
13741  TAATTGGCAA  TAAGATTGAG  ACTATCTTTT  TTTTTTTTTT  GAGACAGAAT  TTTGCTCTGT
13801  GGCCCAGGCT  GGGGTGCAGT  GGCACAATCT  CGGCTCACTG  CAACCTCTGC  CTCCAGGGTT
13861  CTAGCAATTT  TCCTGCCTCA  GCCTCCCCAG  TAGCTGGGAT  TACAGGTGCA  CCACCACGCC
13921  TGGCTAATTT  GTGCATTTTT  AGTAGAGATG  GGGTTTCGCC  ATGTTGGCCA  AACTGGTCTC
13981  GAACTCAGGT  GATCCACCTC  GGCCTCCCAA  AGTGATGAGA  TTACAGGCGT  GAGCCACCGT
14041  GCCCAGAAAA  GACTATCTTA  TTTTATGAAT  TTAAATAATT  GTGAAATTAT  CCACTTAAGG
14101  GAATTAATAA  ATTATAATGT  AATCTTAAAT  TTTAGTTGGC  TTACATAAAG  ACTTAAAATA
14161  CATCAATTTA  AATAAAAACT  CATTTGTCTA  AAAAAAAATC  AAAAATTTTC  CTTGTGCTTT
14221  AAATGTGCTA  CCTCTTTAAG  TTCTAATTAA  GAGAAAAAAA  GTTTAACTGT  GAGTTTCATT
14281  AGTGGTCTTA  GTTAACAGCT  TAAAGTATTT  TGTAAAAAAA  ATACTTCACA  ATTTTTAAAT
14341  AACTTAAAAA  TATTAATACC  TCTTTTATTA  GGTTTTTTTA  ATAAGGAAAA  TATATAATAC
14401  ATCTAATCAA  GATTTTTTTT  GGACAAATTG  GCTTAATAAT  TTCATTTTAA  AAATGGCTTC
14461  TTTATTCTTA  TACTGTAAAA  ATAATATTAG  CAGAATATTA  TAGTATACAC  AAGTTTAGGG
14521  TTCATATTCT  AAAAAACAAA  AACAAAGCT   AATTTAACTT  GCATTTACTA  AATTTCTTCC
14581  ACTAGTTGTA  CTGGTTACAT  GAGTTAACAT  CACTTTATTT  ATTATTCTAA  AATTGTAAAT
14641  TATTCATTGA  ACCAAATTAA  ATGATAATAG  ATAATGTCAT  TTTTAAAAAT  GGAATTAAAT
14701  TTTATGTTAC  TAATTATAAG  GATTCAATGT  GTGAGCTTAA  GTACTGAGTT  CACAGTGTAT
14761  GATAACTTTA  AGAATTTAGG  TGAATATTAT  TAAATTGAGT  AAATTAATTC  TCAATCTTTG
14821  GATACCTGGA  CAATTTCTAA  ATTGGAGGGT  ACAAAATACA  AATCACAAGA  AACAGTGTAG
14881  TTTTATGCAA  ATAACATTTT  TACACAGTTT  AGAATAACCA  TTGATAAACA  GATAAGAGAA
14941  CATATGATTG  CCTTAGAATA  GATACTGTTG  CTTTCGCCAC  TTTAGATTTG  TAAATCACGT
15001  ACTGTATACG  TGTGGGCGTA  GAGGACCATG  CAGGTTTTGG  ATGACTGCCT  CTGTTTTCGT
15061  CATGCCTATG  CGGGAACACA  ATTGCCTGCT  TTGTTTAAGG  GCTATGGTTA  ATCCAAACAG
15121  CTCTGACTCT  ATCAAGTACT  ATAGCTACAG  AGAAACACAA  GTAAGCATTC  GAGATAATGA
15181  CTACCTTGAG  CCTTTACTTA  TTTAAAAAGT  TGTTACTGTT  TGTTAATGTG  GTACATTCAA
15241  TTTACTATGG  ATTGTCACTC  TAAAATAAGA  CTTCAATCTT  TTTCTTATTT  TTATATAGCC
15301  ATGATTTATA  TTCATATCTT  AATGTAATAA  CCAATCTTCT  CTGACAACAT  TATAACAATG
15361  CTGGAACCTC  CATTTTCAGT  ACTTCAAACA  ACAAATACTG  CTTTTATACT  TCAGAGCAGA
15421  TGGATATGTG  CTTCCCAGTG  TAAACACATT  TGGAATCTCA  CTGAGAAATA  CACTATCACT
15481  AAAAATACAG  TTCTGAGATT  CATTAAAAGA  CCTCCAGAAT  TCTGGAAGTA  GGAAGTTTCC
15541  TCTTCAAAGT  CTACAGAGGA  AGATGAGGTC  TGAAATAGAC  AGCTTCTTCC  TTCTTTTACC
15601  TGTGGTATTA  TTCTGTTTTG  TCCTTTTCTC  CATTATCTGT  CTTTCCAGTG  ATGAAATTTT
15661  GATCTGGCCC  TCCCAAGTAT  TAAAAAACAA  GCAAATAAAC  AAATCTCAGT  TATATTTTAC
15721  TAAGATATTG  GCATGCTAAC  TTTTTGCAGG  TTTGTAACAA  GGACCTTTAT  AACTTGACTA
15781  AAAGTTCCTA  AATAAGAATA  TTTACTAGAA  AATTTATTTC  TGCCTGTGGC  CCACATTTGA
15841  GTCAAATAA   TCAATTAGGA  AAAATGAACT  TGTTTAACTA  AAGTTGACCA  AACTGATCTT
15901  TGACCAAACT  GATCTTTGAG  ACCTATTCAT  CTAAGACAAG  CCAATTAAAT  TCTTGGAGAC
15961  AATTTGTACT  TTAAGGAATT  CTTATAATAT  TTGTAATTAC  CCTCATAACT  TTTTTTTTTG
16021  CCCTACTTCT  GTGCTTCTCT  AATATGCAGA  TTATTAAATG  TTGTTACAAA  GCCATTGTCA
16081  AAAAACAAA   AAACAAAAAA  CTAAACAAAC  TCACATGGTT  AGACTTGCTC  CTTTATGAGA
16141  TATTTTTACC  AAAAATGGAG  GAGTTGAAAA  ACTCTGGTGC  CAGAAATCGT  GAAGACATGG
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16201  CCTACCTAAC ATGGAAATGT TGGTTGTCAG TGGAAAATAC TACACAGAGA TAGCCATAGT
16261  GCTGCACAGC CAATCTTAAG TGTTTCTAGA GAATCACTAA TTGTTTCTAG AGAATCACTA
16321  ATTGTTTTCT TTTAACATTC TTGGTTTATA CAAGAAGAGA GTATCCATAC TAAACTCTTT
16381  TCTACTGAAA ATAATGTGCA AACATAACAT CCTATTCCTA GACAGTTTGT AGTTTTTTTC
16441  TCCCATTTCT ATTTTATAAA TCATCTTTTT AAAATACTTT GTTGAGTGAA ATCAGTCCAT
16501  TGCTTGATAT ACCTTGAGCA CAAGTAAATA GTATGCCAAA AATTAAATGT CTTTCAGTCA
16561  CAGTTTGACA AACTCAACTA CCCTGAGCCT ATAGAGTGGT AATAATTGCC CTACTCATAA
16621  AGATGGGGTG AAGATTAAAT GAAATAGCAC CTATAGAACA CTAGTTCCAG ACGTGGTATC
16681  ATGCTAGTAA AATGGCTGCA CAGCACTGCT CAATGATGAC AAAAAGTGAA GCTTCTGGAG
16741  ACAGACTCCA AGTTTGACTC CCAGATCACC ACATATAAGA TGTGGGACTC TGAGGCAGGT
16801  CATTTAATCT CTCTGTGCAT TAGTATCCTT CTCTATACCT TTACAGTGAT GGTAATAGCA
16861  CCTACCTTCT AGAAGTATGT GAAGATTAAA GATCCTTAAT GCATATAAAC CACTGTGTTT
16921  ACTGCTGTTT GACAAATTTT ATTTATAACC ATCTTTACGC TCCTAAAAGG ACTTGAAGCA
16981  GCTTATGACT GAAGACTTTG GTAGGAGTTG GCCTTCTATA AATTATAAGA ATTTCATAAA
17041  TTATTTGATA TGAAAATGCC AGTTGATCAT AGTATGTTTA CCGGGGTCCA ACAGGTTGAG
17101  AAAAAATACA CTTTTTTTCC CTGAACATAT GAAATTAGCT CTCTAGGCAT ATTCCTAAGG
17161  ACTTAAAGAA TGATAACTAT CATTTCTCTT AAATCTTCCA GATTTGGAAG GATATATATA
17221  TTCAGCACAT TGACAGACAA TCCCAGTAGT CCTAAATTAA AAGACATTAA AAATTAGTGA
17281  AACTTTTCCT ACCTTTAGCC TGTGTAATCC TGGATGACCA AGCATAAAAT TAAATTGAGT
17341  AGAGTATACC ACTGTAACAT TTCCTGAAAG GTATTCTAGG CTCTGAGTAA TTTCTTTGGG
17401  GTCTGAAGAT CAGTTTGACA TATCCTCAAG TATCATGAGT TCATTATAAT TAAGAAAAAG
17461  AGAGTAAATC TGGAGAATGA GCCACTTTCT TACTACTCCT TGACCTCAGT TCTTTTTTTC
17521  AGAGACAGGG TCTCACTTTG TTGCCCAGGC TGCCAGGCTG GAGTGTAGTG GCGCAATCGC
17581  ATCTCATTGT AACCTCCACC TTCTGGGCTG AAGCCATCCT CCTGCCTCAG CATCCTGAGT
17641  ATCTGGAACC ACAGCAGGTG CACACCACCA TGCCAAGCTA ATTTTTTAAA AAGTTTTTTG
17701  TAGAGATGGG GTCTTACTAT GTTGCCCAGG CTGGTCTCAA ACTCCTGGGC TTAAGTGATC
17761  CTCCTGCCTC AGCCTCCCAA ATTGTTGGGA TTACTAGTGT GAGTCACTGT ACCCCGCCCC
17821  ACTTCAGTTC TGAGGAGGAA AAAATATGTA ATAATAATGG GACTTTGGTT TGCTGATTTA
17881  AAGATTCATG TAACCTTATC ATCCAATGCG CAATTTGTAG AATAATTAAT AGAGACATCT
17941  GGTCTCATGT TTCTACAGTT GCTCATGCCT TGATAGTAGA TCTCCTTGCT GCTGGCTCAG
18001  AAGGGTAAAA GAGCAGAAAT GATGGGGCTT CTCTCATTCT ATGAGGAAAT AGACCTATGT
18061  AGAGGAGGCT ACCTGTGGTA AAACCTTATC CTCATCACTT AAAATTCTAG GCTTATTCTC
18121  TGACCATATC AAGTTTTCAA ATGGTAAAAG AATTGGATTC AAGAGAAATA TGAATAAACT
18181  TTTGTTTTCA CTTTTCTCCC TCCTCTCCCC CCATTCTCCC TTCCTTTATT TTCTTGTCCT
18241  TAGTTTTCTT TTCACTTTTT TGTCTACTAT TATTTGCCCA AACTCAACTG TAGGCTAGAA
18301  CAAAAAAAAA TTGAAAATTA AAATGTGCCC CTTTTGTTGT TAGACTTGCT TAAACAATTG
18361  GGGTAATGAA CCTTGGACAC TAGATTTTAA AACACACACA TTTGAGCTTC AGTGCACTGA
18421  AATAAATATA TTTTTAACAA TTAAAAAATA AAATTGCATG TTTAAAAAAT CTGCAGAGAA
18481  CAATACACGT TGTGAGATCT TGAATGGAAG GAAAACTGCT AGCCTCAAGA GTGGATCAAA
18541  GATGCTCAGC AGGCAACAGA GTAAGAGCAT GTTGGAGGGT TTAGAGAGTG TGCTCAGGGT
18601  TCTAGGCTCT AAAAATCAGA CAGTCCCCAC GGCCTGGCCT TCGTCGCTGT ATCTTCTTTA
18661  TGAAAAACAC TAAGTCTTTT TCCTCACTGG ATAAATTTTT ATCCTTCAAG TTTAGATCAA
18721  ATGGAACTTT AGGACACTGA CTAGGTTACA TTCATCTTTT AAGAGCGTAC AGACATTCAA
18781  GGGCTAGAGG ATGTGGGTTT ACTGCACAGG CTCATTATCC AACAGCTGTG CTACCTGGGA
18841  AACTTAACCT CTCTGTGCCT TAATTTCCTC ATCTATAACG CAGGGAGAAT GACAGTAGGT
18901  ATCTCATAAG GTTGTTGGAA CAACTAAATG CATTGGTATC TATTGTGTAA AGTGCTTAAA
18961  ACACTGCCTG GCACAGAGCA AACATCCAGT GAACTTTAGC CATCATCATT ATCATTGTTC
19021  TCAGAGTCAA ATACAATATC TCATATCTGA TAAATTACAG AAGTGAATCA ATCACTCTCT
19081  CTCTTTTCTC CAGGGGGAGA CAACAGCTTT TAGACATATC TTTTCCAACA GTCGTCACTG
19141  CTGGACACTG TTTCATCTTG CAAATAAACC AATGAAAATG AGTGATCCTA GAAGAAGATA
19201  AATGGAGGTA TTTTGAACAA TCAAAGAAGG ACAAATGAAC ACCTGGCTGA GAAAAATTAG
19261  CTCTTTTTTC TATGCATAAA ACTATTAAAA TATTCTTCAT AGAAATTTAT GACACAGGAA
19321  ACATAAAGAC AAAATTAAAA TAACTCCTAG TATCTCCTAT TCTTTTTATA TGTATATTAT
19381  ATATACTCAT ATTCATATAT ACATATATCT CACATCATGT ATCATATATA AAATAAATTT
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19441  AGGTGTCATG  ATATATATTT  AGATAAATAT  ACTTAGAAAC  TTTTTTATGG  ATGTATAATT
19501  TATGGATATA  TTGATAATTA  TGTATTTGTT  ATTGACTACT  TCAATTGATT  CCCATTTTTA
19561  TGCATTATAT  TATAGATTAT  ATAGCTCACA  CATCTTTGTA  CATAAATCTT  TGTTCAAATA
19621  TTATTTCCTA  AGGATAGACT  TCATGAAGTG  GAAATACTAA  ATCAAAAGTG  AAAAACATTT
19681  TCTAAGGTTC  TTAACATATA  CATTGCCAAA  TTGCTATTCA  GGATCATACC  AATTTATAAT
19741  CCCAAAATAA  TATGGAAATT  CCTGTTTTAT  AGCACTCATA  TTTACAATAA  ATTTTAAAAA
19801  TCACTGTTAA  CCTAATAGTC  CTTCAAAAGA  AAAAAAAATT  GAAATTACAT  TATTTTAATG
19861  ACTCTATTAG  TGAGGGTCAT  TCTTCCCATG  TTTCTTGTTA  GCCATGACCC  TATAAGAAAT
19921  AAACTGCACT  GCAAAATGAT  AAACATGACA  TCAATCATTA  CATGGGAAGG  CACTATATAA
19981  AGAATAATAC  CTTAGGTTAA  GGCCACATAA  ATATTTATCA  GGTGCCTTTT  CTGCGGAGGA
20041  CTCTGAAGGG  ATACTAAACT  GCATTAGCT   GCATGCAACT  GAAACTACTT  TTACCTACAT
20101  TGTCTCTTAT  AAACATTATA  ACTACTCTTT  GAGAAAGTGT  TTACTATGGA  CTGAATTGTC
20161  TCCCCATCCC  CCCAAATTCA  TATATTGAAG  CCATAAACCC  CAATATGACT  CTATTCCTAG
20221  ACAGGACTTA  TAAGAGGTAA  TTAAGGTTAA  ATGAGGTCAT  TAGGATGGGT  TCCTAACTGG
20281  ATAGGATTGG  TGGCCTTATA  AGAAGAGGAA  GATTCTGCAC  TTGGTCTTCC  AAATTAAATA
20341  ATTTATTTAA  AAGAAAAAAA  AAAAGAGGA   AGAGAGGGAG  CTCTGCACAT  ATACTGAGGA
20401  AAGGCTATGT  GAGCTCTCAC  AGTGAGAAGG  TAGCACTCTA  CAAGCCAGCA  AGAGAGCCCT
20461  CAACAGAATC  CAGCCATGCT  ATACCCTGCT  CTGAGACTTC  CAGCCTCCAG  AACTGTGATA
20521  AAATTTTGTT  GTTTAAACCA  CACAATCTAT  GGTATTTTTT  TATGGCAGCC  CAAGCCAACA
20581  AAGACAGCAT  CATTGCTGTC  ACTTACAGAC  AAGAAAACTA  AGACTAGGAG  AGAGAAAAGT
20641  TAAACTTGTC  CAAGGTCACA  AAAGCCAGAA  ACAAGTGAGG  TGAGAAGTTG  ACCTTGTTCT
20701  CCTCAATCCA  AGGCCAGGAC  TCCTCCACTC  CACATGTAGA  TAGCCACCTC  ACAGTCAACA
20761  GCCAAATGTC  CACACCCCAG  AGTCAGCATT  AGACCAAGAT  GTCTTACCAG  GAGACAAATG
20821  CCTCATCTTG  AATAAATATG  ATCTAACAAC  TTACCCATGT  AAAACATTGA  ATCTCATGAG
20881  AAACAAAAAT  GCAAAGTATG  TAGAAAACTA  TGTTTACCAC  TTAACTGACA  GTGATAAAAA
20941  GCTTAATGAT  ATCCTTATAG  TCTTGGAGGG  GTTTGTATAT  GTGGTGAAAC  AGGTGCTCAC
21001  GCACTGCTGA  TAGACTGTAA  ATTGGTCCTA  GAGAGAAAAA  TAAATAAACT  GGAAGGAGAT
21061  ATGCTGTATG  TTTACTTTTT  TTATGGAAAC  ATATGATATA  CCTGGAAATT  CGATTGACCA
21121  TGCATCTATT  TCTTCAATGG  GTATGCACAG  TTGAGCTGTT  CCCATGCACC  AGGCACTGTA
21181  ATGGGACAAC  TGCACATGAC  AGTCAAAAAT  CTCAGTCTCA  TGAAGTCGAC  ATGCTCATGG
21241  AGAGGTGCTA  CCCACTAAAC  TAATATTTGT  ATATCAATTA  TGGATACATT  GGGCCACATT
21301  TACAGAAATT  CACTTACAGT  GGGTTACCAG  AAGGGATTTT  TTTTCTTGAT  TGGCAAGAAG
21361  GCTAGGCTGT  TTTGTTGGGG  GCTGGCAGGA  GCTGTCTAGG  CTGCCCAAGT  ATGCAGGTCT
21421  CTTCTATCAT  CCTGTGTTAA  CCATCTTCCA  TGTATCTTTC  AACCTCATGG  TCATCTGCAG
21481  CATGTCTAGG  GGTCATATCT  ATGTTCCATG  CAGGAAAAAA  GGGTAAAGGG  AAAGGGAAGT
21541  AGGCATGTAC  CATTTTAATG  CACACCTTGG  TTTTCAGAAA  ATTAAGAAG   AAAGACTTTC
21601  TGCTTTTCTC  TGACTATTCT  GTATTCTGGA  TTACAACGCA  ACAGAAACGT  CACCTTAAAT
21661  TCTAATGTTT  TTCTCTCCTT  GCTTTCAAAA  ACTGACTCAT  TAACCTCCAC  GTGGCTTGGA
21721  AAAATTATTT  CAGTCATCCA  GTAATGAGCT  GTTCATAGAA  ATGTTTGGA   CATCAAGTCT
21781  GTGTTGTTAG  CATTATACAT  GTTAAGCATT  GAATAAAAAA  CAACATGATG  TGGGTAAATT
21841  TCTTTACTTA  CATATAAGTA  CTTATATACT  TATAGCTGAA  AAGAGAGGTT  GAAATGTCAG
21901  GTGGAACAGA  AATAAGATTA  CCTAGATGTT  TCTCCTATGG  GTGATTTTCA  GCTATGCTGA
21961  TCTTTCTTCT  GGGTCAGGTA  CTCCCAGAAC  TTCCTAATTA  AATGGTGGCC  CTGATCTTAG
22021  TTCCTCTCTC  CTCTTAGACA  TTTTCCAGGA  CTACAGAAGA  TGTGCAGTTT  ATAAATGAGT
22081  AGCAGAAACC  TACTGAACAA  ATTATTCAGG  CTCATCTGAA  CAGAGAGGAC  ACCTTCTCTG
22141  CTATACTCTC  TCAGTGATTT  CCCTGCCTTG  GGGTCAATTA  TTGTCTTGGA  CATTGATTTA
22201  AGCACATAAT  AATTGTTGTC  ATTGCTTATG  TTTGGATTTC  ATCTCCCAAA  ATAGATGGTA
22261  AATTCTTTAG  TTTAGAGACC  AAGTAATACT  TAAAAAAAAA  TTTTGTGTGT  GTGTGTGTGT
22321  TTTTTCTGTG  TCTCTCAGCC  CTGTAATAGC  ATCGTACTTA  CACTTGTTAG  ATTTTTAGAG
22381  ACAACTTTTA  CAAAACATGG  AATTATCTAC  ATACCCTTTC  TACAAAACAG  ACAAATTAAA
22441  TACTCAGTAG  TTGAACCAAA  AAAAGCAGTT  CAAATAAAAT  ACTTGAAAAT  GAAGAAATCA
22501  TTTGAACAGA  GTTAAAGTTA  ATCGTAAAAT  AATGTCTGTA  AAAATTATTG  CCAATCAAAT
22561  ATAAAGTTCA  AAAATAGTGC  TTGAAAAAGG  AAGAATCATA  TGAAAAGGGA  CTACTCATTT
22621  TAAAAATGTT  AGATATCAGG  AAAAGCCAAG  AAGTGAGTAT  GGTAAGAGTG  CTGTCAAGTG
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22681  AAACCCTGCT  AATCTCACTG  AACATGTAAA  AATCTGTAGA  TGCCTTTATT  TTATTCACTC
22741  ACACACATAT  GTAGAAAGAG  AAATATATGG  TAAACATTAA  AAAAACCAAA  TTAGAATGTA
22801  AAATTAATAC  TTTAAAAAAT  GGGCTGTATA  CTTTTCTTAT  CACCGGAGAT  AAGAATTTAT
22861  TATTTTTAAA  ATAAAGTTAT  TTTCTCTGTG  ACTGTTCCA   TGACTTTGCT  ACTTAGAAGT
22921  TAGAGATGCC  AAAGTTTATC  TAAGAAAATG  TTTATGGAAA  TATTATTTCA  ATAATGAATG
22981  TTTAGAAGAC  TGAATTTCCT  GACTGGGCGC  AGTGGCTCAT  GCCTGTAATC  CCAGCACTTT
23041  GAGAGGCTGA  AGAAGGAGGA  TCGCTTGAGT  CCGGGAGTTC  AAGAGCATCC  TGGGCAACAC
23101  AGCGAGACCC  TGCAGCAAAG  TAAAAGAAA   AAAGAATTGA  AAAGGAAGA   CTGAATTTCC
23161  TTTGGGCAAG  TCATGTGACA  TTCCTGTGCC  TCAGTTTCTT  CATCTATAAA  GTTAATTCCT
23221  ACATTTTTGG  GGAAGGGAGA  GAAAAACTTA  GGATAGTGAC  TGGCACAGAA  GAAGCACTAT
23281  ATACTATATA  TATGTGGATA  TCATTTGTTT  TTATGGTACC  ATTTTAGCTA  TCTAATGCAA
23341  AATATGAATC  TTTTTTTTCT  GGGTCTTAAA  TTATGGAATG  TAAGAATTTT  CTAAATTCTC
23401  TAATTCTGTG  TTAGTTTTAA  AGCAATGGAG  TAACGTATCT  GTCAACTTGT  AAATATAAGG
23461  ATCAACCTGA  TCCACAATTT  GACCCCTAGC  CACTAATATT  TAATAGTACA  ACACTCAGAA
23521  ATTATCAAAG  GTCAGAGAAG  CCAAACAAAT  GTAAAAACAT  ACAGGTGCTC  AGAAAGATGC
23581  ACCTGTAATC  TCTCTAAGGA  GAAATATTTT  CCAAACTGAG  TGACACGGTG  CTTTAGTGAG
23641  TTGTGGAATC  AATCTCATGA  TTTCCAACCT  AGTGTTCTTT  TAAAAATGAA  CTAGTCCACA
23701  GTAGAATATA  CTAAAGTGCT  GGTGCTTAAG  ATAGTATTGT  TTTCTGGAAA  AAAAAAAAAA
23761  ATTTTTTTTT  TTTGAGACAG  GGTCTCGCTC  TTGCCCAGGC  TGAAGTGCAG  TGGCACAATC
23821  ATGCTCACTG  CAGCCTTGAC  CTCCTGGGCC  CAAGTGATTC  TCCCACCTCA  GCCTTTTGAG
23881  TAACTGGGAC  CACAGGTACG  TGCCACCACA  CCCGGGTAAT  TTTTTAATTG  TAGAGACAGG
23941  GTCTTGCTAT  GTGCTTAGGC  TGGCCTTGTG  AACTCCTGGG  CTCTAGTGAT  CCACTAGCCT
24001  CAGCCTCCCA  AATTTATGGG  ATTATAGGCA  TGAGCCACCC  TACCTGGCCT  GTTCCCTGAA
24061  TTTTTTTTTC  TTTCAGGTGT  TTGTGCATAT  GTGTGTGTGT  ATGGGTATAA  CAGAGAGACA
24121  GAGAGAAAGA  AACTTTTCTA  TCTCACTTTG  CAATCAGAAG  TTTGAAGTCT  TATCTTTTGG
24181  CTTTTGTTTC  AGAAATATTT  CAAATGTAGA  CTCTCTCCTT  TACCACACTG  TCCCCTTAGG
24241  CAAGGTCTTT  GCCATTCTTC  TGAGACTATT  GCAACAGACT  CCCAACTTCT  GACTGTGGGC
24301  CCTTCTCAAA  AATGATTGTT  TATGCAATAA  ATCTAAACCC  AAGACAACTA  CAACAATACA
24361  ACAAATTCTC  TGCTTAAAAA  CTTCCAATGT  CTGCCGGGCG  CGGCGGCTCA  CGCATGTATT
24421  CCCAGCACTT  TGGAGGCAGA  GGCGGGCAGA  TCACTTGAGG  TGGGGAGTTC  GAGACTAGCC
24481  TGGCCAACAT  GATGAAACCC  CATCTCTACT  AAAAATACAA  AAAATTAGCC  AGGCATGGTG
24541  GTGGGCGCCT  ATAATCCCAG  CTAATTGGGA  GGCTGAGGCA  GGAGAATTGC  CTGAACCTGG
24601  GAGGTGGAGG  TTGCACTGAG  CCAAGATCAC  ACCATTGCAC  TCCAGCCTGG  GCAACAAGAG
24661  CAAAACTCTG  TCTCAAACCA  AACCAAAACA  AAACTTCTAA  TATCTACCAA  ATGTTTCACA
24721  CAAGTATTTG  GGGATCTTCA  CAAATGGCCC  TTATGGAGTT  TTCCTTTGCT  GAGACCCTAT
24781  GCTCTGGCCA  CACTAAACTC  ATTCAGCATC  CCAGAAAGGC  CTCAGCCTTT  GTGAGCAAGC
24841  TCTTATCTCC  AGGCCTCTCA  CAAAGACCTG  TTCCAGTAGA  AGCTCAGGGG  AGCACACTGG
24901  ACATTATTCC  AACAACCCTT  TCCCCACAGC  TATGCAGCCA  AATCTGCCAG  CTCAGTTAAT
24961  TAATTAAGCA  ATTCAGAGAT  GAGGGTCTGC  CCAGGCTGGA  GTGCAGTAGC  TGCGACCTCA
25021  AGCTCCTGGG  CTCTAAGTGA  TCCTCTTCAG  TCTACCCAGA  AGCTGGGACT  GCAGGCATGT
25081  GCCACCACAC  CCAGCTAATT  TTTTTTTTTT  TCAGTAGGGA  CCAGGCCAAC  CTAGTCTTGA
25141  ACTCCTGGCC  TCCAGCCTTC  CGAAGTGCTG  TAATTACAGG  CATGAATCAC  TGCGCCCAGC
25201  CAACCCGCCC  AGTCTTGTTA  GACATGGGGT  CTGTAGTTTC  TAGTAGGTTC  TTGAGTCTAG
25261  GGTTCCTACC  TCATGTTTTA  TAGTTAATTT  AGGGGAGGGA  CTGTGTCTGT  TTATCTGGGG
25321  ATGTAGGGGT  GGGCAGGGGG  ATAGAGGGGA  CTTCAATTAA  TGAAACCAGA  AGCAAAACTC
25381  AGTTGAGGAC  ACCGGTCATG  AGAGTGGCCT  GATTATGGCC  AATCTTACAT  AATGTGTGAG
25441  ATCTTGATAT  TACCCCATCC  TTGAGAGTCC  TCTATAAAGC  TACAGGGACT  TGGGAGCACC
25501  TTTAATTACA  GACAACCCAT  GTTCCTGTGG  ATTATGATTT  ATTAGATTGC  ACATGCCTAA
25561  ATAAAGACAT  CCTCTGCAGT  CTTTTGACAA  TTCTATAAGC  ATCTTCTGAC  TCCGCAATTA
25621  GACAGCTAAG  AGATCTGTGT  TACTTCCCTC  ACATATATAA  ATAATTTTAA  ATAAAAATCA
25681  TGGCGTGAAT  AATTTCTTTC  CTCTACCGAT  TTGAAGCTAT  CCATTTGGAA  GACCACTCTG
25741  AAGAGATGAA  ATAAGTCTTC  TGCCAAAGAT  TACTTATTAA  TTTACAAGGA  AAAGGGGAAG
25801  TTTTGTTCCT  CTCCGTGAAT  TTGATTGAAA  ATCGAGGGCT  TTCTCGAATA  GTTTTGGCAT
25861  CCAGGGTCAT  TTTTCATTAA  AAAGAGAAAA  GTCATGTCAA  ATATGAATTT  CCGCAGATTA
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25921  TTCAGCACTA GACCCTGGGA GATTCTGTAA AGAGGGGTTT TGTTATACTC AACTTTTCCG
25981  GGTAAAACAA ACACAAATAC TCCTCCTCCA AGGGGCGGGG GCGGTGCCTA GGTGATGCAC
26041  CAATCACAGC GCGCCCTACC CTATATAAGG CCCCGAGGCC GCCCGGGTGT TTCATGCTTT
26101  TCGCTGGTTA TTACATCTTG CGTTTCTCTG TTGTTATGTC TGAAACCGTG CCTGCAGCTT
26161  CTGCCAGTGC TGGTGTAGCC GCTATGGAGA AACTTCCAAC CAAGAAGCGA GGGAGGAAGC
26221  CGGCTGGCTT GATAAGTGCA AGTCGCAAAG TGCCGAACCT CTCTGTGTCC AAGTTGATCA
26281  CCGAGGCCCT TTCAGTGTCA CAGGAACGAG TAGGTATGTC TTTGGTTGCG CTCAAGAAGG
26341  CATTGGCCGC TGCTGGCTAC GACGTAGAGA AGAATAACAG CCGCATCAAA CTGTCCCTCA
26401  AGAGCTTAGT GAACAAGGGA ATCCTGGTGC AAACCAGGGG TACTGGTGCT TCCGGTTCCT
26461  TTAAGCTTAG TAAGAAGGTG ATTCCTAAAT CTACCAGAAG CAAGGCTAAA AAGTCAGTTT
26521  CTGCCAAGAC CAAGAAGCTG GTTTTATCCA GGGACTCCAA GTCACCAAAG ACTGCTAAAA
26581  CCAATAAGAG AGCCAAGAAG CCGAGAGCGA CAACTCCTAA AACTGTTAGG AGCGGGAGAA
26641  AGGCTAAAGG AGCCAAGGGT AAGCAACAGC AGAAGAGCCC AGTGAAGGCA AGGGCTTCGA
26701  AGTCAAAATT GACCCAACAT CATGAAGTTA ATGTTAGAAA GGCCACATCT AAGAAGTAAA
26761  GAGCTTTCCG GGAGGCCAAT TTGGAAAGAA CCCAAAGGCT CTTTTAAGAG CCACCCACAT
26821  TATTTTAAGA TGGCGTAACA CTGGAAACAA GTTTCTGTGA CAGTTATCTA TAGGTTTAAG
26881  TTGTGATGCA GCTGAGTTGA AAAGGCTTGA GATTGGAGAA TTAATTCAGG CCAGGCTTCA
26941  AGACCATCCT GGGCAACATA GCCAGACTAC CATCTATACC AGGGGTCCTC ATTTCCCCGG
27001  CCACCGACCG GTAACCGGTC CCTGTCCATG GCACGTTATG AATTGAGCCG CACAGCTGAG
27061  GGGTGAGCGA ACATTAACCA ACTGAGCTCC ACCGCCTGTC AGGTTAGCTG CAGCATTAGA
27121  TAGATTCTCA TAAGCTCAAA CTGTATTGTG AATGGCACAT GCAAGGGATC TAGGTTTCAG
27181  GCTCCTTGTG ACAATCTAAT GCCTGATGAT CTGAGGTTGG AGCAGTTTTA GTCCGGAAAT
27241  CATTGCTCCC AGCCCCTGCA CCCCCTGGTC CGTGGTATAA TTGTCTTACA CAAAACGGTC
27301  TCTTGTGTCA AAAAGGTTGG AGACTACTGG TTTTACAAAA AAGTAAATTA GTCAAGCATG
27361  GTTGGCACGC TCCCTTAGTC CCTGCACCCA GGCGTTTAAG GATACAGTGA GCTATGATGG
27421  TGCTACCTCA CTCCAGCCTG GGTGACAGCG AGTCAGACGT TGTCTCAAAA CTTAAAAAAA
27481  AAAAAGTTA AAACAGAAAA AGGGCTTCTT GTCAGAGACT GCCGTATATC TAGAGGTCCA
27541  GGAACTAAAA AGTCTGATGT CCAATCCTGA AAAGCTCGAT GGTGCACTAG AGGAGGCTTT
27601  TACATGTAAG AGCATCTAAG TTCTGGAAAT GCCAGTGTCA GGGAAGGGAA GTGGAGAGCA
27661  ATTTGGCATC CAAACATAAC TTGCTGATAC TTTTTTTTTT TTTAACACAA GTACTACATT
27721  CTAGTCTTTC TGTGGTGTCA TTGTAACTAT TGTTTCTTAA TATGCTATCC ACTGACTTCA
27781  AGGGATCAAT AAATAGGAAT CAAGGTGTCC CAGAATATGG ATTAGGGGAG TTTTTTTGTT
27841  GTTGTTGTTG TTGTTGTTTT TCATCTATTC ATTATCCTGT AGCTGAAATT TAGAATTTTC
27901  TTCCATTGTG TGTGACTGAT AGAAATAACA AATTTGTAGG TTATAGTTGT TGCAAGAATC
27961  TGGAAATCGT GCTTGCTTAT TTCCGAAGTA CTATTAGGTA TATCAACAAA AACACACATA
28021  TTACGGTCAA GTGGTTTGAT AATTATTTTA ATATTATTGG TCTAATACAA TTGTAACCCT
28081  ATGAATTACT TTAAGTATCT TATTTATGAA AAGAATCTGT AAGTTTCATC AGACTACCAG
28141  AGCATACCGA AGACTGAAAA ATTTTAAGAA TCCAAACCTT AATGGAAATG TTGGAGGCTG
28201  CCCAATTAGG TTCTGAATTC CACCTTCCTG AATCACAAAC TTGTTTTAAC TCTCAGTCTG
28261  AGGTAAACTA CGTTTCTCTT TAAACAGACA TAGTTTAATT TTCCTTTGAT TTTTGATTTA
28321  GTATTCTTAC TGATCATCAT AAATAACCAA TGCTAATGTT AGTCTACTTT GGACCATGGT
28381  ATTTCGAGAA ACTTTGAACA AAGTCCCCTG CAAAACTATG CATTGCATTA TTTCACATAC
28441  ATTTATGTTT TCCAGACGGT TCAATAGTAC CTCACTTTTC TGAACTTATT TGTATAGTTT
28501  GGCATCTTTT TAAAAATTGT GTCCTATAAT GAAAGGTTGT AAACATTATG TTTTAAATTT
28561  GTATAGATAA AATCAACCAC AGACCTTTCC TTGCTTGGAT GTAATTGCCA TTGTTTCCCA
28621  ATGAGTTCGG AATTACTAGG ATTGTGCAAA AATATGCCTC ACTTGCCTGA CATAGCAGAG
28681  AGCCATTTTG CCTAAATGCT GTGCCCAGCA ATGGACTGTC ACCAGATTCT CATCACATAC
28741  AGTGAGGATG AACAACTAGC CTCTCCCAGC AGCTGGCCGG TCTCTCAATA ATATGGGACT
28801  CCCTCAAGAT GGCTTCCTGC ACCTTTGCTC CTCTAGCCTT GTATGTATAC AAGGCTAGCA
28861  TGCCTGGCAT ACATAAGGTT AAAAACAAAA TCAATAAGTT ATGGTTCTTC CTCCAGTTCT
28921  GGGGATTATT AGACCACTTT TTTGTTTTGT TTTGTTTTGG ATGGAGCCTC GCTCTGTCAC
28981  CCAGGCTAGA GTGCAGTGGC ACAATCTCGG TTCACTGCAA CCTCTGCCTC CTGGGTTCAA
29041  GCAGTTCTCT GGCTCAGCCT CCCACGTAGC TGGGATTACA GGTGCCGCC ACCACGCCCG
29101  GCTAATTTTT GTATTTTTAG TAGACGGGGT TTCACCATCT TGGCCAGGCT GGTCTTGAAC
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29161   GCCAGACCTC GTGATCCACC CACCTTGGCC TACCAAACTG CTGGGAATAC AGGCGTGAGC
29221   CACCGCGCCC GGACTTAGAC CACTTTGTTT TGGCCAATAG GACAACAGCC ATAGAACCCT
29281   CCGCAAATGA GAGCTTGTCC CTAAAGATGC TTTATTTACA TAGCTGTGTG CCGCATGAGC
29341   CAAAAGGTGA TAACCTTTGT TCAACACGCG CCTCCAGCCC TTCGGTTAAG TCCAAAGTAC
29401   CATTCTTAGA ATGCTCTAAA ATACATAATT TTTTTTTTTT TTTTTTTTT TTTTTTTGAG
29461   GAGTCTCTCT CTGTCTCCCA GGCTGGAGGG GAGTGGCGCG ATCTCGGCTC ACTGCAATCT
29521   CTGCTTCCGG GCTAGCTGGG CCTACAGGTG CAGACCACCA CGCCCGGCTA AGTTTTGTAT
29581   TTTTTTTGGT AGAGGGGGTT TCACCATTTT GGCCAGGCTG GTCTCGGATT CTTGATCTCA
29641   AGTGATACAC TAGCTTTGGC CTCCCAAAGT GCTGGGATTA CAGTCGTGAG CCACTGCGCC
29701   CAGCAAAATG CTTTTTGTGG AGCCAATCAC TTTATTAGCG CTTACCTCTC TATGCCTACT
29761   TTATGCTTTG AAATTTTGTC ACAGTGTGGC CGGTCATGGC AAACACAATT CATTCTTATG
29821   CAGGATGTCA CGGTTATTTC TGTCATCCAA ACTCATTCTC GCAACGCATT TCAGCTCTTT
29881   AAACGACTTT GTGAGCGGCC CTGAAAAGGG CCTTTGGGTT TTTTTGTTTT TGTTTTTTGA
29941   AGTTCTCAGG AGACCGCGTA TTCTTAGATT CAGCCGCCGA AGCCATACAG AGTGCGCCCC
30001   TGACGTTTTA GGGCATATAC TACATCCATG GCTGTGACAG TTTTGCGCTT GGCGTGCTCC
30061   GTATAGGTGA CGGCGTCTCG AATAACGTTC TCTAAGAAAA CCTTAAGCAC ACCTCGAGTC
30121   TCCTCATAGA TAAGACCGGA AATGCGCTTG ACGCCACCGC GCCGAGCCAA ACGGCGAATA
30181   GCCGGTTTTG TAATGCCCTG GATGTTATCC CGGAGCACCT TACGATGGCG CTTAGCACCA
30241   CCCTTCCCCA AGCCTTTTCC GCCTTTGCCG CGACCAGACA TGATTCCTAT CGCAGTGGAA
30301   GGTATGAACT GAAACAGTTC CTTAAATACA AACTTGGCGG ACCTGATTGA AAACAACATG
30361   AGTTGGCGCG GTTTTTTTTT TTTTCAAAT TTGGTCACCA AGTGGGTGGA GCAAGAAAAA
30421   CTGTTTCATT ATGGTTCATT GTTTTGATTG GCCAGTGACA GCTTGCTCTT TGTGGGAGTG
30481   GAAGGGTGTT TGCAAGTTGA ATGCGCTGTA TTCCTGTCAG CTTAATGACG CTAAGCATAG
30541   CCCCATTCCA CATTTCTTTT TATTTCCACT TGCTAACTAA TAAATTACGG AATAGTTTAT
30601   TGGGGAACAT ACAAATAATG TTTAAGGAG GTCAGATTTA TAGGTCAAGG GATTTACCCT
30661   CCCAATCATT TTAATATTTT TATTTAAACC AGGCATTTTG ATGGCCTTCT CTGTGCTGGA
30721   CAAGGTATAA GTTTGGCTAT GAAGTTTCAC TCCTAAAGAC CCTATGTTTT GGGAAGGCAA
30781   AAAGGTAGCC AAATAATTGC AAATTAAAAC CTCATAAGTG CAAACTTCTT CCTCGTCACT
30841   TTCCCTATCT CGATTCAAAT ATTTGTTGAA TGACTCATTT TTCTGCAAAA GTCTGAGAGA
30901   GACAGGGAAT ATAAACTTAA GTCTGGATAA TATGTTTTCC CGGGACGCTC TTCCTGGTCT
30961   GCTGTGCCTG TTTGCTGTGC CTGAAATTCC AAACACTCTT CCCTTCCCTC CGTTTTTAAT
31021   CCCCTTTCAA CTTGCTACAG CTTTAGAGAA AAGAACATTC GTTTTGTACA GTTGGGGATT
31081   AATTGAAGTG TAGGGCTAAT ACTTGATTAA GGTCATTACA AAATCTACAG GGTCTTCCTC
31141   TGGGAGGTTT TTGTGATAAG ATTATTGGTG TTAAAATAAG GCTAATCCCC TTGAAAAATA
31201   AATAGAATAG CAGAATTGGG TCTGAATGTG GTTTGAAGAA AGGGACTTCT CAATTCAAAA
31261   TTTTATTCTT AGCTTCCTGC GGGAGCTTTC CAGAATGCCC ATAAGATCCA CTTTTGTTTA
31321   AAAAACAAAA ACAACCCCAC CCACCACTCT CTGGTTAATA AATGAATTTC TATTGGGAAT
31381   ATTTAGAATG GGGCTGTGGC CTGTGAGAGA CATTATATAG TAACCTCAGA CTTGCTCACA
31441   TGAAGAGAAG AAATCCAGGA ATGGAGAAAA AAGACCCAGG AAAGGCCAGA ATGCTCTACA
31501   TGTCATATTG TTTGTATCAC TTCTGAAATA ATTGATTACA TTCTTCTGCC CCAAATTGAG
31561   TTCTTAGGTT CTTCCACTCA CTGTCCACAT GCCACAACAC AGACCTTATA ACTAGAGACT
31621   TAGCTAGGAA GAAATGTCAA ACATTACAGA GAAAAAATGC AGAGTCTGAG ATCATAAGTA
31681   AAACTCTGAA ATCTCAACAT GCCTTTTAAT TCATGAAAAT AAAAAATATA GCAGCATATG
31741   CAATATGACA ATTCTCTGAA AACATACATC ATGTGAACTA CCCTGGAACA CATCTCGCCA
31801   AGTGCCATCT TCATTTTAAC CAGAGGTCTA GGATGCCTTT CCTTTATTTT GCCTATTATA
31861   TCATTTATAA AACCCCATTT TTATTTTGAT ATTTTATTTA CTTTCTATTT CCTGCTCCTA
31921   ATATCTCCTT TCTAAACTTT TCTCAATGAC AGTGACTCAA AAACAATGAA TGTCAGAACA
31981   AATATTTAAA GGATCTGTAC ATGTAGATAT ATATATTTAA AATGGATTCT TCCACTCTGC
32041   GAAGAATTCA GGCATACTCA ATCTTATGGT TAGGGAGAGA TTAGGCTCAC TCGCCTAATC
32101   TGTATGGCTT CTCGTTCGCT TTCCATTTCA CCTTCCTCTC ACCCATCAGA TCAAACTCAT
32161   TCATTGAACA AGAGACCTAA GCCCTTCAGA TTAAAACTCT GCAAACAAGT GTGTGGTTGAG
32221   AGGATACATG AAGCATTCAA ACAAATAAAT CTATGATATT AATCAGAGGT TAATCTATGA
32281   TATTAATCAG AGGTTAATGC AGTGGCTCAC GGCTGTAATC CCAGCACTTC AGGAGGCTGA
32341   GTTGGGAGAA TCGCTTGAGC TCAGGAGTTC AAGACCATTT TGGGCAACAT AGCAAGTCTT
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32401   CATCTCTACT TAAAAAAAAA TAACCAGAGG TGTTATGAAA ATATAAATTG TCCAGAACTA
32461   CCCTCCACAA ACTAACTCTC TCAGAATATT CGATATGAGG AATGAAATAT GGTGTGTGTG
32521   TGTGTGTGTG TGTGTGTATG TGTGTGTGTG TGTGTGTGTA TGCACCTATA TATGGCACCT
32581   ATATATTCAA CAAACAATTC TGATAATTGG CCAGGGTTGA GAATGACTAG CAGCCCAGCA
32641   TACACTATCA GTTTTAAGTA TATAATTGCG CTTTAGTAAA ATGTAAAGAA ATCCCAGAGT
32701   AGAAATACTT TTAAGCTATA TTACAGGTGA GAAAATGCAT AAGTATAGTC TCACCCAACT
32761   TAGACTATGG GGGCTTTATA ATGTCACAAC AGTTGTTTCC AGGCATTTGG GGACATCACC
32821   ACTGGTCTTG GGCAAGAAAC TCCTCTAGCC AATGGCTGAT TTATCTCACT CCCATCTAAG
32881   GCTTCACTGC ATTTCTCTTT TTCAGCAACC TAACTTATTT AAAAATATCC ATTTTCTGAT
32941   TCATTTTTTT CTGAATTAAA CTGTCAGTAC CATTGGCACA CCTTTGGTTC CGTAGCATAC
33001   CTGTGTCTCT GCTGTGTTTT TTTTTTACCT CCACTCCTTA CTTTTCTAGA AAAAAATCTC
33061   TGCTTTTTCT TTTCAGTTTA AATTATTTCA CAAAAAGTTT TCTTGACTTG CACTTCCTAG
33121   GCTTGCTGTC CTTGTGTGGG CACGCTCCCA TAAACACTAT TAATACACTT CGATTTGTTA
33181   AAAATAAAGA TATCTGGACA GAAAATTTCT TTTCTTTTTT TAAGATTTTA AAATTTTTAA
33241   TGTTTATTTT TTTCCTAGAC TGGAGTACAG TGGCACCATG ATGGCTCATG GTAGCCTACA
33301   CTTCCCCGGG CTCAAGTGAT CCTCCCACCT CAGCCTCCCA AGTAGCTGGG ACTACAGGTG
33361   TGCACAACCA CACCTGACTA ATTTTGTTTA TTTGTTTGTT TTGTTTTTTG AGATGGAGTT
33421   TCGCTCTTGT TGCCCAGGCT GGAGTGCAAT GGCGGGATCT CGGCTCACCG CAACCTCTAC
33481   CTCCCAGGTT CAAGCAATTC TCCTGCCTCA GCCTCCCGAG TAGCTGGGAT TACAGGCATG
33541   CATCACCACG CCCAGCTAAT TTTGTATTTT TAGTAGAGAC GGGGTTTCTC CATGTTGAGG
33601   CTGGTCTGGA ACTCCTGACC TCAGGTGATC TGCCCGCCTC GGCCTCCCAA AGTGCTGGGA
33661   TTACAGGCGT GAGCCACCAC GCTCGGCCAC TAATTTTGTA TATTTTGTAG AGATGGGCTT
33721   TCCCTGTGTT GTCCAGGCTG GTCTTGAATT CCTGGGCTTA AGTGATCTGC CCACCTTGTC
33781   CTCCCAAAAT GCTAGGATTA CTGGCGTGAG CCACCAGGTC TGGCTGGAAA GATAATTTCT
33841   AACATTATCC TCTCTTAAAC ATTTGTTTCA AAAATTTTAC AAACATGAGA GTAATTAAAT
33901   TTGATTTTCA AAATTCCCTT GAATACTTTC TTAATAGCAC ACAGAAAGCA CAAAGTATTT
33961   TACATTTGTT TTAATGATGA AATTGTGAAC CCAAACTTAC ACAAAGAAAA ACCCGTAACA
34021   TTATACCCAT ACTTAAAACA GATGCCCTCA TATACATAGT AAAACTCTTG GGGGCAGTAG
34081   TGAAGTTGGT TATTTACTGT TTTATGAAAG TGCCATTCAG CCGGGTGCAG TGGCTCATGA
34141   CTGTAATCCC AGCACTTTGG GAGGTCGAGG CAGGCTGATC ACGAGGTCAG GAGTTCAAGA
34201   CCAGCCTGAC CAAAATGATG AAACCCTGTC TCTACTAAAA ATACAAACAT TAGCTGGGCG
34261   TGGTGGTGTG TGCCTGTAGT CCCAGCTACT CAGGAGGCTG GGGCAGGAGA ATCGCTTGAA
34321   CCTGGGAGGC GGAGATTGCA GTGAGCCGAG ATCGCACCAC CGCACTCCAG CCTGGGAGAC
34381   AGGGCGAGCT CCGTCTCGAA AAAAAAAAAC AAAAAGTGC CGTCATAGTG ACTCAGTTTT
34441   AAGGAATAAA TCAAGGATAT TTAACTCAAT AGACTACAGT TAGCTAACGT GACTTGCACT
34501   GAAAGTTATA CGAATATTGG TACTTATTCC CCTGCCCCTG AAGTATGAAT TAAAGACTCC
34561   AAAATTCTTT TTAGAATCTT CAGAGTAAAA GCTAGAATTT GATTTTTTA AATAATAAAA
34621   AAATACTTTG TATCTAAATC TGGTGTATAA AATAACTTGG TGGATGATGC TTCAAGGCTA
34681   TCCATCCCCA AATTTCTCCC TGAATGATAA AGAGAATAAA TGAATATGTC AATTCAAAAG
34741   TTAGAAATTT GGCCGGGCAC GGTGGCTCAC TCCTGATAAT CCTTTCGGAC GCTGAGGTGG
34801   GTGGATCGCA TGAGCTCCGG AGTTCAAGAC CAACCTGGGC AACATAGCCA GAACCCGTTT
34861   CAATAAATAA TAGAAAAAAA TGAGCCAGGC GTGGTGGTCC CAGCTACTCA GTAGGCTGAG
34921   GTGGGAGGAT CACTTGAGCT CAGGAGGTCG AGACTGCAGT GAGCCGTGAT CGCAGTACTG
34981   CACACCAGCC TTGGTGTCAG ACTGAGACCC TGTCTCAACA ACAACAAAAC AAGTTAGAAA
35041   TTTGGCTGGG CGCGGTAGCT CACGCCTGTA ATCCCAGCAC TTTGGGAGGC CAAAAAGGGC
35101   GGATCATTTG AGGTCAGGAG TTCGAGACCA GCCTGGCCAA CATGGTGAAA CTCCATCTCT
35161   ACTAAAAATA CAAAAAAAT TAGCCGTGCA TGGTGGCATG CGCCTGTAGT CTCAGCCACT
35221   TGGGAGGCTG AGGCAGGAAA ATTGCTTGAA CCCAGGAGGC AGAGGTTGCA GTGAGCCGAG
35281   ATCATGCCAC TGCATTCCAG CCTGGGTGAT AGAGTGAGAC TCCATCTCGA GAAAAAAAAA
35341   AAAATTCTGT ATGAACTGAA CAAAATATCC TTAAATTTTA AAATACATCT GAAAGATATT
35401   TCAAATATT TAGGAAAAAA ATTATAGGGA TCAGGCAAAT TCTGAGATTC CTTTTTCCCT
35461   GCAGCAAACA TTAGGAGTGC TGCTGTTCCT AAAAACATGG TAACTGTTGC CACACCGTAT
35521   GTTTCCTTGG CTCAGACATA AGGTTGTGTA GTTGTTATTC CAGAATAGCT AGAATAAAAA
35581   TCCAGCACAT CATTTTCTTC AGCAAGTTAA CTAACCTCTC TGTGCCTTGG TTTCATAACA
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35641   GCAACATAAG   CATAACAGAA   TAGCAGCAAT   AGCTCCTACC   TACCTCATAA   GATTCTTTGG
35701   AGGAATTAAA   TTAAGATTCA   GAACACAGCC   TAATATCTAG   TAAGTAATAA   TAATTGGCTA
35761   AAAAAATTTT   CTTAAGATTA   TATATATTCA   TGGGGTACAA   GTACAATTTT   GCTACATTAA
35821   TATATTGCAT   TGTGGTGAAA   TCAGGGCCTT   CAATCCATCC   CGGAAAAAAA   AAGTTTTTGA
35881   AAAGATTTCT   GCCATGGAAA   ACTTTTAATG   TACAAATTCA   TCCATCCAAG   AAATAGAAAA
35941   TATATAAGTA   TCAACTCCAA   ATCCACCATA   TCTATCTCTT   CTACACCTTA   AACAATTACT
36001   CAGAAATAGA   ATGCTTGAGA   TACCAGAATG   CATGCATATC   AAGTAATAAA   TGCATGCAGG
36061   ATGTCAACGC   ATCCTAGGCT   TTCAAATAAA   ATTGTCATAC   AAAATACTTT   AATATTGTAG
36121   TAACATTCTA   CATGTTAGAG   TGTAGAAGTT   AATCGCTGAT   GCAAAAAAGG   AAAAGAACAC
36181   ATTATACCCA   AAGCCTACAG   AGAGAATCAC   AATTACAAAT   ATCAGCCTGC   ATGTGAAAAT
36241   CTTTAATTTG   AAAGTCAGAA   ATATTTAAAT   GATAGTCATT   GTTAAATCAG   ATTGTGGTTT
36301   GAAAAAAGT   TAGTTTAAAA   CTGAGTTTAT   GAAAAATTTG   GGGATTTTAG   AGACAGTGTT
36361   TTGTTTTTAA   ATGTGTGTGA   GTTTGTGAAG   AATGTTTTAT   AAAATACTGA   CAGTATTATA
36421   AGATGACATT   ATTATAATAC   AACATAAGAA   TTTTGGCCTG   TACCTCTCAG   CAGTCCTCAA
36481   TCACCTGCTG   TACTTGACTC   AATGATTATC   AGAGTGGTTT   GTTTTCCTTC   TGTTGTGTTC
36541   CCAGTTCAGG   CAGCTCAGCA   ATGGCCTGTG   ATTCCAGCAA   TTCAAATAGC   TGGTAAGTAG
36601   TTTCTTGTTT   GTTTTCTCAA   ATTTTCAGGG   GCTTTTCTCT   ACAAGTGATT   TCCAGTGCAC
36661   GCCCCTCCAC   CCATTCTTTA   TTCCTTTACC   TTCAGGAAAA   CCCTCAGCGC   TGCATCTCTG
36721   GTCACCGGAC   CACCGTGGTA   CATTTACCTA   TGGCCACCAG   GTGTCACCCT   TCTCTTTACT
36781   ACCATGGTTT   GTGAATGGTT   TTGCCAGAGG   TGAATAAGAA   TTTAAAATGC   AGGTCTTTGA
36841   TTTTTCAAAT   GTAGTTGACC   TTAAGAATTT   ATGAATAAAG   CCAGAAAAAT   TAAGCTTAAA
36901   AAACACCGAA   AGAAAATGAG   GACTTAAAAT   TTCTATTAAA   AAAATTAACA   GGCCACAGTT
36961   GCTGATGTTT   AGTAAATGTG   TTAGTGAAAT   GTGTTACTGT   GAAGACTGGG   GTGTTTCTTG
37021   AAATCTCAGC   CCAGGTGAAA   TAAAACCAAT   ATAAACAAA   TGCTTACCTA   ATAAATTAAT
37081   TGTAACATAT   TCCTTATGAG   GTAGAAGAGT   AAGTGAAGCC   TTATAGCAGT   CTGCTTTCAG
37141   TATAGTAAGA   TATTAAGAGA   GAAATAATTT   GTCATATGCT   TTCAGAATGG   TTTGCTGGTA
37201   AAATAACCAA   TGTCTTACAA   CTTAGACGAC   AATGTCCCTA   GAGTGAAGAA   ACACGATTAA
37261   TTCGGCTACC   ACAGTTGAAT   GAAAATATTC   CGTAAGACAA   AATGTAAAGA   AATTAGAAGC
37321   AAAATAAATG   TCTCCAAAAT   GACAAAGCGA   TTAAGTATAT   ACACAAGATG   AACAAGAACT
37381   TCAATAAAAT   CATGCAGTAT   ACAATACAAT   ATACATTTAT   TAAAGTATAT   GCATTTTTAA
37441   TGCAACAATA   ATACTAACAG   GTAATAGACA   AGTTGTTAAT   AGTTTTTCAC   TGGCTAATTA
37501   AATAACAGCT   TTAATTGTAT   TCATTTATA   GCTTTTCTAC   AATGAGCGTA   AATCACATTT
37561   ACTTTTTTCT   ACATAACTTT   TCTAACCACA   AAAAAAGAAA   ATGGTTTAAA   AGAAGAGATG
37621   AGATATCTTT   GCTAAAATTT   AATGCCTAAA   GAAGAAACTT   CTGAGCTGTA   TATGGTATCC
37681   TGAAGCACCT   GCCCTTCAAG   ACAGAATGCT   TGTACCACAT   TTATGCAGCC   AAGTGCATGT
37741   AGTAACATAA   AGTAAACACA   TGCCATCTGG   ATATATATAT   TAAGACTCTT   TTGACGGCTG
37801   GGCAGGGTGG   CTCACACCTG   TAATCTCAGC   ACTTTGGGAG   GCCGAGGCAG   GCGGATCACG
37861   AGGTCAGGAG   AGTTCGAGAC   CAGCCTGGCC   AACATGGTGA   AACCCTGTCT   CTACTAAAAA
37921   TACAAAAATT   AGCCGGGCAT   GGTGGTGCAC   GCCTGTAATC   CCAGCTACTT   GGGAGGCTGA
37981   GACAGGAGAA   TCGCTTGAAC   CTGGGAGGCA   GAGGTTACAG   TGAGCCGAGA   TCATGCCATT
38041   GCACTCCAGC   CTGGGCAATA   GAGTCTCAAA   AAAAAAAAAA   AGACTCTTTT   GAACATGGTG
38101   AACTGATTTC   CCAGAATCTA   GCAATTCCTG   AATGTCCTGG   TTAGATTTTT   TTTTAATGT
38161   GCACCGGAAC   CCCAGTGGCT   CCATGGAAGG   ACCTGGGCAT   CCTCTAAGCC   ACTTGGTGGC
38221   TTCCATTATA   CCATCTCAAA   ATGAGAGAGC   TTACTCCACT   TCATTGAGGG   AAATACCACC
38281   AGAGTTCTGA   CTCCAGAGGC   ACTGGCCTAG   GGAGGACACC   GTGTGTGAAG   CCCAGCAGGG
38341   CCACTAGCTG   TCCCCACCAA   TTACAGTCCT   TGCGTAGGGT   CCAAAGAAAT   GAATGCCAAA
38401   GAGAGCAACA   GAGGAGCAAG   GGAGTCACAT   TCCAGGACCT   TCCTTCAGGG   ACTTTAAAG
38461   GAAACATGAC   AGCTGAGGAT   CAGTTGGTTG   TTTTCTGCTG   TTCCCCTTCA   TGTGATTCAA
38521   GCTCACTCAG   AAGAAACACA   ATGAGACAAG   AGAAGAGCCA   TCTCCTTCCT   TCTCTATTTA
38581   TTCTAGGCAT   CTAAACTACT   GAATGTAGTG   GTGTCTGAGA   TGTATCAAAC   GGTCAGATTG
38641   ACTGAGTTTG   AAACCTGTTT   CTATCACTGA   CAAACTATGA   GATACTCTAT   ACTTCACTTT
38701   CTTTTTTTT   TCATTTTTTT   ATTTTTATTT   TTATTTTTTT   GAGATGGAGT   CTCACTCTGT
38761   CACCTAGGCT   GGAGTGCAGT   GGCGCAAACT   CGGCTCACTG   CAAGCTCTGC   CTCCTGGGTT
38821   CATGCCATTC   TCCTGCCTCA   GCCTTCCGAG   TAGCTGGGAC   TACAGGCGTC   TGCCACCACG
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38881   CCCAGCTAAT TTTTTGTATT TTTATTAGAG ATGGGGTTTC ACCATGTTAG CCAGGATGGT
38941   CTCGATCTCC TGACCTCGTG ATCCACCCGC TTTGGCCTCC CAAAGTGCTG GGATTACAGG
39001   CGTGAGCCAC CGTGCCCGGC CTACTTCACT TTCTTCATTT AAAAAGAAA TGGGGATAAT
39061   AGTACCTATC TCATAGAATT ATTGTAAGAA GTGCATGCAG TAATGCATGT AAGTAGGTGC
39121   TCAGAAGAGT CGGACACGAA GTAAGTGCTT TTATCATCCT TATCATAATT TTCATTATCA
39181   GAACAAGGAG AGACCAGGTA GAAAATTATT GTGATTCTTC AGGTCTGGAA TACTAGAGTA
39241   GCATCCCAAA TGAAGGCACC ATTAAACTTT GCAAATCTGT ATGACACCTT CATGCCAATT
39301   AGAAAAAACA CCTCTTCACA ACCCCTTTCA AGATATTTGC CTCCTACCTG CTAAAAACAC
39361   CCATCATACT ACCCACAGAT AGCCATGATG CTTTTTCTGG GACAGGTGCC TCTTCCATTC
39421   GTGCAGTGTA CAGCCTTCAT AGCTGTGCAA CTCACATCAC AATCAGATGG AAGAATCCCC
39481   AAGGCTTGGT GACAGATGAG TTACTGGGTA ACACAGAGAG AGGATTCAAA GGAAAGTTG
39541   AACGGGTCCA GAAAATGCAT AGATACATGT GTAAAAATCT GGTAAGGTTA TGACTAGCCA
39601   CGTCCCAGGG TTCAAAGCTT TTCTCAGATG TTAAAATGAA TCATGTAAGT CCCCCAAATT
39661   TAAGGAGTCC TCTTCCAAAA ATAGGAAATG AAATGACATA GGTGTATGTC TCTGAGGTGA
39721   CGGAGGAAAT GAAGGAAGCC TCTAGATGCA GCTTGAGGTT CATGAGAGAC AGTTCCAGGG
39781   GAGAGGTCAC AGCTAGGGAT CACCGGCATG CAGGAACTCA GAAACCTAAA TGGGGAAATC
39841   TTTTTGAGGA AATGAACAGA GAAGGCTAAA ATCAAGGAGT TCGTCAGGCA ATTTCTATGT
39901   TTAGGTTCAA CTCTCTCCTG AAACATGAAG AGCTCATAAA TGCACTCCCT CTTTGAGTCT
39961   CTAGTTTTGT CTCCTTCCCA CAGTGAGTCT GCAGGCTGCG TGTCACTCAC GTTCAGCTAA
40021   GACGTAGTGC CCCATGGCTC CTCCTGTGGA GACAAGAGAC CCAGGAAAGA GGCATCACAA
40081   ACCTAGGCAC CATCTTGCCT CTTCTCTCTT CCTTATTTTC CTCATTCACC CATCTCAATT
40141   TAGACCTGGG CACTATTGGA TTTCAAGAAC CATTATCTCT CATCTGGAAA TGCTTATTGG
40201   CTTTCTAACT GGTCTCCTCA CCTCTCATCT AACTTCTTAA CAACACATTC ACCATATAAG
40261   GGAGATCGTG GTCCTCCTTT CTTAGGATCC TTCAATGACA CCCCAGTGAT CATAACCCAA
40321   TATCCCAAAA GACCCTTGGA CTCTGTATGA GCTGGCTTCT TTCTGATTCT CTTTTCCCTA
40381   CACCACAGAT GTTCAGGGGG TAGAAATGCA TAATTGGTGA GTGATAGCTA CGCAAACTCA
40441   GGGTTAAGGT ACAGTAATTA TTTCTAATCT CCCAGTATGC CTTATACTCT CCTACTTGGC
40501   ATGGTTGCTC CGTCTGTGTA GACCTCCCAT CATCTTCAAC CTCACCTAAT GGAATCCAGC
40561   TTCTCCTTCA AGATCCAGAA GGCTATCTTG ATCCCCAGCT GAATGTGATC ATTCTTTCCT
40621   TTGACACCCT AAGCATTTGC TTCCTGCCTG CTTTAGGACC TCATGGGGTC TTCTTTAACT
40681   ACATTTACTT GCTATCAATT TCATTCCCTA CCAGATTTGG GTTCTGAGAA TAGCCACAGT
40741   GACTTCTCAA CCTCAAAGCC CCTGTACTAC CTTAAACAGC TCTTGCAAAA TAGTAGGTGC
40801   TCTGAAGATG TTTGTTGAAT TAGAGACTTT CATTCTGGGG AGAACCATTA TTTTCTGTCT
40861   CCCAGGGAGC TGCTGGTGTC CCCAAAGAAT ATAAATGAGA AAAATGCTTC CCATGGATGC
40921   CAGATCCCCT CTGCCCCTCT TCCCACTGTG CCCTGGGGCA GAGGTACTAA GAGACTTCCC
40981   CCTTGTTCCT ACTCACTTGA ACCCTGCCTC TTCCTTAATA TTATGAACAA AATTCCAATG
41041   AACAAGATGA CGACAAAAAC AGCAATTCCA CTGATGACTC CAATGACTAG GGTGCCAGAC
41101   GGTGAGGGCT CTAAAACAGA AAAAGCAAGT TAAAGCCTTT GATTGCCACC CTCAGCCCAC
41161   CCCCTAACAA AGAGCAGATC CTCATCTCAC TGCCATAATT ACCTCCTCAG GCACTCCTCT
41221   CAACCCCCAA TAGATTTTCT CAGCTCCTGG CTCTCATCAG TCACATACCC CAGATCACAA
41281   TGAGGGGCTG ATCCAGGCCT GGGTGCTCCA CCTGGCACGT ATATCTCTGC TCTTCCCCAG
41341   GGGTACAGC CAAGGTTATC CAGCCCTGGT AGGTCCCATC CCCATTGGGC AATACGTCTT
41401   TAGGTTCGAA CTCCTTGGCA TCCATTGGCT GCTTATCCTT CAGCCACTTC ATGGTGATGT
41461   TCTGGGGGTA GTAGTTCAAG GCCCGACACC GTAGAGTGGT CACTGAAGAG GTCACATGAT
41521   GTGTCACCTT CACCAAAGGA GGCACTTGAC AGGAAAGAGG AAGGATGAGG AGAGGGGATC
41581   TGTTACCCT TGCCAGGAAG ACTGGAACTT TCACTTCCTT CTATAGGTTG GAGGAAGGAA
41641   ATACCCTTTT CAGAAAAAAA CAAGCTACAG GAGAGACACC ATTTTGTGTC CTAAGATTGG
41701   ACTCTAACAC AGTGTCACTT GGAGAGCAGT CAGATCAGCT TGTTCTCCTC ACATGTAAAT
41761   ATACATATCT GTTACCCATG TTCTTTGTTC TGATAGATAA AATTGCCCTT TATGTGCATT
41821   GAAAATGATT GAATACAGAT GGTCAGTTTC ACCTGGGTCA ACCTAGGAGG CATTGTTATA
41881   AGAAGCGGAC TTGTAAGATA GGTAGCTTCA GTGATTATTG CTATGTTCTA TGAAAGAAAC
41941   TTTTAACCTA AAGGATTCTT CTACTCTGAT AAGTGGCCTC ACTTGATATT TTGTCCTGGT
42001   ATTCATATGA TAGCTGAGAT CTCTGAATTC TCTTTTTTTT TTTTTTTTTT TTTTTAAGAT
42061   GGAGTCTCAC TCTGCTGCCT AGGCTGGAGT GCAGTGGCGC GATCTTGGCT CAGTGCAACT
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42121  TCCGCTTCCC AGGTTCAAGC GATGCTCCTG CCTCAGCCTT CCAATTAGCT GGGACTACAG
42181  GTGCGCATGA CTGTGACCAG CTAATTTTTG TATTTTTTTA GAGACGGGTT TCACCATGTT
42241  GGTCAGGCTG GTCTCAAACT CCTGACCTTG TGACCACCCG CCTCGGCCTC CCAAAGTGCT
42301  GGGATTACAG GGGTGAGCCA CCGTGCCCGG CCTTGACATT TCTGAATTTT TAACAGGTAT
42361  AAATATACAA AAGATTATTG GTTAAATAAA AAGCAAGGGC CATAGACACT TCCCTTTGAG
42421  CCATATGCAT GGAGAAAAGA AATTAAACCC ATGACTTGTG GCTGTCTCAT ACATCTCAAT
42481  TATAAGGTAG AGACTCTAGG ATTGAGAAAG TCCCTTCCCA GAATTTGGAG AGGCACACAG
42541  CCTCAGCCAC CTCTGAAACT CCAACCAGGG ATTCCGTGCC CTGCAACCTC CTCCACTCTG
42601  CCACTAGAGT ATAGGGGCAG AAGTGTGTTT CCACCATACC TTGTTGGTCC AAAACACCTC
42661  TCCCCAGCTC CAGCAACTGC TGCAGCTGTG CAGGGCAGTC CCTCTCCAGG TAGGCCCTGT
42721  TCTGCCTGGC CCGAATCTTG TGCCTTTCCC ACTCCAGCTT GGTGGGCCAG GCCCTGGGTT
42781  CTGCTGCTCT CCAATCCAGT GTGTCAGGGC AGAATTCAAG GTGGTCCTGC CCATCATACC
42841  CGTACTTCCA GTAGCCCTCG GTACTGTTGT CTTCTTGCAT TTCACAGCCC AGGATGACCT
42901  GCAGGGTGTG GGACTCTGGA AAAATCCCCA GCCTTGTTAA CTGCAACCAA AGGAATAGGT
42961  CCCTATTTCC ACCATCCCCA AGGACCAAAT GATCTCAGGA AGCAAATTCC TTCCCTCTTC
43021  CCTGCTCCCA CAAGACCTCA GACTTCCAGC TGTTTCCTTC AAGATGCATG AAAAGATGAA
43081  AAGCTCTGAC AACCTCAGGA AGGTGAGGCC CCCTCTCCAC ATACCCTTGC TGTGGTTGTG
43141  ATTTTCCATA ATAGTCCAGA AGTCAACAGT GAACATGTGA TCCCACCCTT TCAGACTCTG
43201  ACTCAGCTGC AGCCACATCT GGCTTGAAAT TCTACTGGAA ACCCATGGAG TTCGGGGCTC
43261  CACACGGCGA CTCTCATGAT CATAGAACAC GAACAGCTGG TCATCCACGT AGCCCAAAGC
43321  TTCAAACAAG GAAAGACCAA GGTCCTGCTC TGAGGCACCC ATGAAGAGGT AGTGCAGAGA
43381  GTGTGAACCT GGAGACAGAG CAACAGGCCT TAACCATGTG TAGTAGGAGG GGAGCAGGAT
43441  GTTGAGGCTC CACACACCTG CATCAACTCA TACCATCAGC TGTGTCTGGT CCTCATTTTG
43501  TGAAGGGTGA GTTGCAGTCC TGTCTTTCTT CCATATGACA GTCCTGGGTG CTCTTTCCTT
43561  GTGTGCTTTT CTCTGCCACA CGTGGCTGCC ACCCCCTCAC TGCCCCCAGA TCCTATTCCA
43621  ATACTCATGA TTAGACAGAC TCCACTAAAG CTGGTGGATT CTAGAAAATG TTAAGGTGTG
43681  TCTAGCCATG GTAGTTGAAC TCAGGAGTTG GTGCTCAGGG CAAATTAGAC CCAAATCCTG
43741  AGGAATAATT CCTTCAGTTT TTTTTTTTTT TTTTTTTTTT TTTTTGAGA CAGAGTCTCA
43801  CTCTATCACC CAGGCTGGAG TGCAGTGGCA CAATCTCAGC TCACTGCAAC CTGCACCTCC
43861  TGGGTTCAAG GGATTCTCCT ACCTAAGCCT CCTGAAAACC TGGGACTATA GGCGTGCGCC
43921  ACCACACCAG CTAATTTTT GTATTTTTAG TAGACATGGG GTTTCACCAT GTTGGCCAAG
43981  CTTGTCTCAA ACTCCTGACC TCAAATGATC TACCTGCCTC AGCCACCAAA GTGCTGGGAT
44041  TACAGAAGTG AGCCACCGTG CCCAGCCTTG GTCCTGAATT CTTACACTGA ACTGCCTATG
44101  TGGCCTCACC ACTTGGAAGC CTGACTGGAA TCTCAAACTT AACATGTCCA AATGCAGATC
44161  CTTGATTTAC CCCAAACTGC TCTTTCCTCT GCCTTCACCA TCTCAGAAAT GGCATTGCCA
44221  ATTACCCCAC TGCTCAGGCC AATAAAATTA AAATAAAGAA CAAAGTCAAC TTTAACTCTT
44281  CTCTTTTTCA GGGGGTCAGG GGAGACAGGG TCTTGCTCTG TCACCTAGGC TGAAGTACAG
44341  TGGCACAGTC ATGGCTCACT GCAGCCTCAA CTTCCTGGGC TCAAGCAATA CCCTCCACCT
44401  CAGCCTCCCG AGTAGCTAGG ATCACAGGTG CATGCCACCA CACCCAGCTA ATTTTTGTAT
44461  TTTTTGTAGA GAAGGGGTTT TGCTGTGTTG CCCAGGCTGG TCTTGAACTC CTGAGCTCAG
44521  GAATCTGCTC TCCTTGGCCT CCTCCTTGGC ATGAGCTACT ACACCAGCC AATTCTTCTC
44581  TTTCTCTCAC ACAACATAGA ATCCTTCAGC AACTTCCTTC AGAATATATT CAGGAGACAA
44641  TGGTTTGTCA CTCCCTTTTC TGTTCCACC CAGCCCACTC CACTACCTCT TGCCTGGACT
44701  GTGTAACAGC TTCCTGGCTG GGCTCCCTGC TTTTACTGTT GCTCCCTTCA TTCTGCTTTC
44761  CACATAGCAG CCAGAGCAAT CTTTTAAAAG CCTGTGACAG ATCACTGTTA CTCCTTGGCT
44821  AGAATTCACA CCACAGCCTA CAGGCGCCTG CACAACCTTG TTTGTGGCTC CTCTTCTGAG
44881  CCCATTACCT ACTTCTTGGC CTCTACTCCC CAGCACTACT TGTTTATTTT TTTCAACCCG
44941  AGCTTCTTAA CCAGGAGTTT GTCTACTAGG TGACATGTGG CAAAGTTTAG AGACATTTTT
45001  GGTTGTCAAG ACTGGGGGAG TGCTCCTAGC ACCTAGTGAG TAGGGAGGAC AGGATACTGC
45061  TAGACATCCT ACATGCAGAT GGTAGTCCCC CTTCCCACCC CCACGCCGCC CCCCCCCCCC
45121  ACACACACAC ACATGAGTAG TGCTGAGAAA ACCCGCTTTT TAATCCAACT TGCCAGGCCC
45181  ACTCAGTTTG CCTGGGAAAT ACTGCTCCCA GTCAATATCA TTCTTATTTC CTTCATGTCT
45241  CTGCTCAAGT GTCAGCCCCA GAGTGACTTG CCCTGACTTC TCTGCTTCTC ACAACACCCA
45301  TGATTTCCTG ATGTTGTATA TCTTTCTGCT CATTTGCTTA TTGTCATCTC TCCCACTAGA
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45361   ATGCAAAATA  TCAAAGGGTA  AAGACTTGTT  TCCCTGCTCT  CTCCCTTGGG  GCTTGAACAG
45421   TGCAACACAT  GGCTGGGACT  CATTTACACT  TGTAAACAAT  GAATATTTCT  GCTCAACATG
45481   AAATTTTATT  ATTCAACCTC  TAATGCAGTG  TGATGTTTAA  GAATCATAGC  TATGAAGTGG
45541   AGACATGAGC  TCTGCCACCA  AAGCCCAGTG  TACCATTGAA  TAAATTTGCC  AGGAAGCAGG
45601   CCGTGCCATG  CCTCATTCTT  GTCATGTGTA  AAATGTGGAT  ACACGTAGTA  CCAAAACTCA
45661   AAGTGCTGTG  CTGAGGCCGG  CGTGTGACCC  ACAGAACACT  GTGCTACACT  ACAGGGCAAA
45721   ATCACTGTCA  ACTAAGATTA  GAAGCAGCTG  TAGTACTTGA  AATAACATCA  GAAAACCAGA
45781   TTATTTATGT  TCTTTGTAAC  CTGAAAAGAG  TTATATAATC  TGAATTCCAG  TTAACTTCTA
45841   GTAAATAAA   CGTATTATTA  GCTCCTACCT  CCCTATGCCT  AGTGAAAATC  AAATAAGATC
45901   AGATATGAAT  GTAACTTAGA  AGTGAGTGCA  TTGCTTACAT  GTTCATTATC  AGTACTTTGT
45961   AGAGAGGCCT  CTTAATTACA  CAGCACATTG  CAAATCAATA  AAGCCTAGCC  GAAAAGAGAA
46021   TTGTTCAGTT  CAAACGTTCA  AAACTAACAT  ATACTTAATT  TTCCAGGCAA  AAGAACAATT
46081   GCCAAGAGTG  GGGAAAGGCC  CGAGGTAGGC  CTCTCTCAGG  AGCCTCCCAC  CCTAGAGACC
46141   TCCACCCCAG  GTCTCACCAA  AAGTGGGTGG  AATGGTGAAG  AATTCAGATC  CCCAACGCCA
46201   CTCTTTCGCG  CCCCCACCGC  CCAACGCATT  CGTTCTGAGG  TGGAAACCCC  GTGCGGATCC
46261   TGCTGTGGGT  TTGCTCAGCC  TTCTCGGCAA  GCACTCAGGG  AAGAACTTCC  TGTTTGGAGA
46321   TGACTGGGGA  AAAAACTGCA  CAGCTGACAT  TGGAAATAAA  CCCGAGTTCC  AGGTTCAAGG
46381   AGCCCCAGGC  TTAGCTCAGC  TCAAGTGAGG  AACTACGAGA  TTTATTTAAA  AGCATTCTAG
46441   TTGGGGGAAG  GGAGTGGGCG  GTTCCAAAAG  TCACTCCGCA  GAGCCGGGAC  AGCCGGGGGA
46501   GGGGGCAGGT  CCTGGGGCGA  GGGACCCCTA  TCTGCAGTTC  AGTGGTAGGC  ACTCCCTCAC
46561   GGGGTCTGGA  CGCAGAAAGT  AGGGAGAGGG  GCTTGCGGAT  AGGGTTGAGC  AGGTCCTCCA
46621   AAGTTAGCAA  ACTCCCAAGC  GCAAAGAAAA  AGCTAGTTTC  GATTTTCCA   CCCCCGCCGC
46681   GCCCCTAGTT  CGCCCGCAGC  CCTCGGACTC  ACGCAGCAAG  CGCCCCTGCA  GGACCGCGGT
46741   CTGCAAAAGC  ATCAGGAGGA  GAAGCGCCGG  CCTGGCTCGC  GGGCCCATTT  CCCCAGCTCT
46801   GGCCGCACGT  CCCCGTTAAA  TCTCCGCTTC  TTTTGGGGGG  CGGGGAAACG  GGGATGGCTC
46861   CAGAAGTCAC  CCTACAGCTA  TTGCCTAGGC  TCAGGAGATG  CCCAGTAAAA  CTTCCTGGTG
46921   AAAAGCAACA  GGTCTTTCAG  AACTTTAGTT  CTCTCTCTCC  TACAGCAGAA  GGTACCTGCT
46981   TGTGAAACAC  TAGGTGATCC  AGTGTCCCCC  TTGGTTTTTA  AATCCTGAAG  GGGTGTTGTT
47041   GATTGGGGAA  AGTAGCTTCG  CAATGTTCTG  ATCTGAACTT  TAGATATTTA  AATATTTATG
47101   ATTTTCAAAA  TTCAATCATA  CATTTAAAAA  TTTTATCTCA  ACCTTAGACC  AACTTATGTC
47161   TTATTTGACT  TAGAAATATA  AAGCTTTTTC  ATTTTGTTTT  TTGATTCAAA  TTAATTAAGT
47221   CATAACATTA  ACCAATTAGA  TCCTACTGAA  ACACCTTCCA  CAGCCTTCAT  AATTGAATTA
47281   TCTGACAAGT  GTTTCACAAA  CTTTACAGTA  TTGGGATTAT  CTGGAGAATG  ATTAAACATA
47341   TTGAGGCCTG  CTCCTAACCC  CAGACACACT  GATTTAATGG  GTAATTGTTA  GGTAGTTAGA
47401   CATTAGCAGT  TGGGAGGGGA  TGACAGAAGA  GAGCGGAAAG  GCTGTCACTA  AGACAGCCAC
47461   TGGCCCACCT  AAATTCAGGC  CCAAGACTAC  CCTAATGCCA  CCCTAAGGGA  TGGAGTTTAT
47521   GATAAAGTCT  GTGGCCAAAA  TATCCTGGAG  AAAGAGAAAG  GAGGGTACAG  GTGGAAATTC
47581   CCTAAGGTGG  CACATGCCCA  ACAACACAAA  AGCCTGTCTT  CAAGTTCACC  CCAAGTTCAT
47641   CATGCCATCA  TTATAATAGA  ATTTACATAC  AGTTTTGCCC  CCCCATCCCT  GGGAGGCTTT
47701   TCTTAACAAA  TTATAGGTAA  GACCATGCAC  AGTTTAATTT  TAGATTGTAT  AGCTATACAC
47761   TTCAATCAAA  TAACATCATC  CTGTCACTCA  GATACAGCCC  AAACCTCAAC  TCCTCCCCAC
47821   AAACCCCATA  AAAGCACCTT  GAGCTCTGTA  AAGAAGTGCT  GAGTTCACTT  CGCAGAAATA
47881   AGCCCGCTGT  CCCTCAGAGT  GTATTATTGT  GCTTCAATAA  ACTTTGCTTT  AAGCTTGCAT
47941   TTTGGTGTTA  GTTTGTAGTT  CTTTGCTCAC  TATCACAAGA  ACTGAGATTG  CTGGTTCAGA
48001   GCTCCGGCTA  TAATAATCTC  CTCGGTTAAA  GGATCCATCC  CAATGCATAA  TTCCCAGTAA
48061   CAGTATGGGA  TGCCACCTGG  GCAATGGGAT  TTTAAAAGCT  TTCCTTCTCC  CTCAACGAAG
48121   TTTGGGAATT  ATTGCCTTAG  ACATTTCAAA  CAATATTAAT  AAATTTAATA  CACCTGATTT
48181   GCTCCAAACC  TTTACATATC  TAGCAAATTC  AACAGGCATT  ATTTTTGTAA  GCATGTATGC
48241   AAATTTTGGC  AATTCAAGAA  AATCAAACAG  GATATCAGGG  CCTCGACTGT  AGGCAAACAG
48301   ATACAATAAC  ATTGGAAACA  TGTAGAATAT  TGATGATGGG  CACATTGGGG  CTGATAGTAC
48361   TATTCCTTTT  TTTCAATTTT  TGGTAAGATA  TAATTAGCAT  ACCATATAAT  TCATCTATGT
48421   AAAATGCAAA  AATTGGCCCG  GCTCAGTGGC  TCACGCTTGT  AATCCCAGCA  CTTTGGGCGG
48481   CCGAGGAAGG  CAGATCACCT  GAGATCAGGG  GTTCGAGACC  AGCCTGGCCA  ACATGGTGAA
48541   ACCCCGTCTT  TACTAAAAAT  ACAAAAATTA  GCCGGGCGTG  ATAGCAGGCA  ACTGTAATCC
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48601  CAGCTACATT AGAGGCTGAG GCAGGAGAAT CGCTTGAACC CGGGAGGCGT AGGTTGCAGT
48661  GAGCTAAGAT CGTGCCATCA CACTCCAGCA TGGGAGACAA GAGCAAGACT TCATCTCAAA
48721  AAAAAAAAAT TAGCTGGGTG TGGTGGCATG CACCTGTAAT TCCAGCTACT CGGGAAGCTG
48781  AGACAGGAGA ATCGCTTGAA CCTGGGAGGC GGAGGTTGTG GTGAGCCGAG ATCATGCCAT
48841  TGCACTCCAG CCTGGGCAAC AAGAGCGAAA CTCCGTCTCA AAAATAAAAT AAATAAAATA
48901  AAATGCAAAA ATTAATGGAT TTTAGTATAT TTACAGAGAT GTGCAACCAT TACCAAAATT
48961  TTACATTTCT ATCTCCCCAA AAAGAAACCA TGTTCCCCTA ATTCAGTACC CTTAATTCAT
49021  CGCCTCCCAG ATTCCTCCAT TCTCCTCCTC CTCCCCTCCC AGCCCTAGAC AATCTTTAAT
49081  CTACTTTCTT TCTATTTGGA ACATTAGTA TACATAGAGG CATATAATAT ATTGCTTTGC
49141  CGTGACTGGC TTCTTTCATT TAGCATAATG TTTTTATGTA TGTTTTTCAT GGACCAATAA
49201  TATCTATTAT AAGGACATAC CACAACATAT TTTATTTATT CATTCATCAG CCGATGGACA
49261  TTGGTTTGTT TCTACTTTAT GGCTATTGGG AATAGTGCTG TTATAAACAT TTATGTACAA
49321  GTTTTTTTGT AGACTTATGT TTTGATTTCT TTTGGTTATA TATCTAGAAG TGGGTTTGCT
49381  GGGTCATATG GTAACACTGT TTAACCTTTT GAGGAATTGC CACATTCTTT TCCAAAGTAA
49441  GCATTTATC CTCCTATCAG CAGTGTATGA GAGTTCTGAT TTCTCTCCAT CTTTGCCTGG
49501  GTTTTTGAAT CAGGGCCCCA GATAGAACAA AAATGTGGTT ATTCAGTTGT TCCACCATCA
49561  CTTGTTGAGA AGACTCTTTT TTCATTGAAG TGTTTTGGCA CCCTTATCAA AAATCAATCT
49621  ACCATAAATG TGAGAGTTTA TTTCTGGAGT CTCAATTTTA TCCCATTATG CTATAATCTA
49681  TAATCCTATC TTTTTTTTTT TTTGACAGAG CCTCACTCTA TTGCCCAGGT TGGAGTGCAG
49741  TGGCCCAATC CCGGCCACTG GCTCCTCCTC CCAGGTTCAA GCAATTCTCC TGCCTCAGCC
49801  TCCCAAGCAG CTGGGATTAC AGGTACCTGC CACCATGCCT GGTTAATTTT TGTATTTTTA
49861  GTAGAGACGG GGTTTCACCA TGTTGGTCAG CTGGTCTGG AACTCCTGAC CTCAGGTGAT
49921  CTGCCCACCT CAGCCTCCCA AAGTGCTGGG ATTACAGGCA TGAGCCACCA CACCCAGACT
49981  ATAATCCTAT CTTTATGTCA GGACTACACT GTCTTGATTA CTATAGCTTT TTAGTAAATT
50041  GAATTCAAGA AGTTTCTCAA CTTCAAATTT GATCTTTTTT TGGAAGACTA TATTAGCTAT
50101  TCTCAGTCTG CTGAATTTCC CTAGGAATTT TAGGATCTAT TATCAATGTC TATTCTATTT
50161  TTGTATATGT TTTAATATTT TCATAAGAAA CTTTTTTCAT TTAAACTTTT TTTTTTAAGA
50221  AAAATAGTGA AAATCAGAAC ACTGGGGGTC AGGCGCATTT AACAGGCAGA AGAAGAATAA
50281  AAACTTGTCA TATAAACAAA AAAGAAATGA CCAATCACAT TGTGGAAGCC ATGGAGTGGT
50341  TATAGGTGCC AAAGGCTGCA GAGAAATGGT GTCAGATATA CCTGAAAATT GTCCATTGTA
50401  TTTGGCCATT AAGAGACTTA GAAGACTTAA GCCATAGATT GCTCAGTGAG ACCCCGAGGG
50461  CAAATGGTCT GAAGGTGAAT AGATCATTTC ACCTTTAAGA GAGCAGGTAG GAAGCTATAA
50521  ATCCAAGATT AAAAAGTTGA CTGAACTGTT AAGGAAGAAA CTCTAATCTT GAGCCACCCT
50581  ATCCTGGCTC CACCTTCTGC TGCAAGCAAA CAGAAATGCT GAAATTCAAC ACTCACAAAG
50641  GCTGGTAAGC TGGAAATGAC AAAAATTACT CCTGGGAAAG TCAGATTTAG AATTAGGCCA
50701  TATTTGTTGG GGTTCAGATT TTCATGTACA CTTGGGAAAG GGTTTAGCTT ATAGGCACAT
50761  GCATGAAGGG AACTGGTATA GGGCTGTGTT CATAAGGTCA AGAGTTGAAG GCCAGGCATG
50821  GAGGCTCTTG CCTGTAATCC CAGCACTTTG GGAGGCCGAG GCAGGAGGAT GGCTTGAGCC
50881  CAGGAATTCA AGACCAGCCT GGGAAACATA GGGAGATGCT GTCTTCACAA AACAATTAAA
50941  AAATAAAATT AGTCAGGTGT GGTGGCACAC ACTTGTGGTC CCAGCCACTC AGGAGGTTGG
51001  GAAGATCACT TAAGCCTGGG ACATTGAGGC TGTAGTCAGC CATGATAGTG CTACTGCACA
51061  CCAGTCTAGG TGACAGAATG AGACCCTGTC TCCAAAAAAA GAGCTGTATC CACATCCCAG
51121  GAAAGTGGTT GAAGATCTAC TTTTCTCTGT AAACCTAATA AAGAATAGAG TGACAAATGT
51181  GTGTTGTGGA AAGAAATGGG GTGAGAGCTA CGTAGATGCA AAACAATACA TCCCCACATA
51241  CCACTTGTTA ATCATCCTTT TCCACCCACT TATGGGATGA ATTGCATCTC CCCAAAAGAT
51301  ACTCTGTCCT AACCCTCAGT AGCTGTGAAC CTGACCTTAT CTGGAATACG GTGAGTTCAC
51361  TGGTTAAGAA GAGATTATAG TGGAATAGGG TGAGTCCTCC AACCAATGAC TGGGGTCCTC
51421  ACAGACACAG AGGGATGATG GCCAGGTAGA GATGGAGGCA GAGATTGGAG TTATGCTGCC
51481  ACAAACCAAA CACAGGAAGC TGCTAGAAGT GGAAACAGGC AAGAAAGAAT CCTTCCCCAG
51541  AGGCTACAGA GGGATCTTGG CCCTGATAAT ACCTTGATCT CAACTGGCCT ACGTAACTGT
51601  GAGAGAATAA ATTTCTTTTG TTCTAAGCCA CCCAGTTGAT AGTACTTTGT TACGGCAGCC
51661  CTAAGGAACT TGATATACAT TTCTTTTACT GTCATAGAAG TTTTGAATCT TTTAAGTAGG
51721  TCTGTACCCT TCCTCCCAGT GTCAACACAT GGAATTCCTC TCCTTGTGCC TTGAAAAGTG
51781  AAAGGTGTTT GAACTGGTAA TGAAAGAAAT CTCAGCATGA GGCCAGATGC TGTACCTCAC
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51841   ACCTGTAATC  TCAGCACTTC  GGGAGGATGA  GGCGGGCAGA  TCACTTGAGG  TCAGGAGTTC
51901   TAGACTACTC  TGGCCAACAT  GGTGAAACCC  CATCTCTACT  AAAAACAAAA  AATGTTATCC
51961   TAGCCGGGCA  TGGTGCCTGT  AGTCCCAGCT  ACTCAGGAGG  CTGAGGCAGG  AGAATTGCTT
52021   GAACCCGGGA  GGTGGAGGTT  GCAGTGAACT  GAGATCACGC  CACTGCACTC  TAGCCTTGGT
52081   GAGAGAGCAA  GACTGGTCT  TAAAAAAGAG  AAAAGAAAAA  TGAAATTTCA  GCATTATAGA
52141   ATAAAAATGT  TTCCCCTTCC  CCCCAAACTT  TAAAAAAGCA  GAAGTCTGCA  TCATAAAATG
52201   GTCTTTGCCA  ATGTTATTTT  TATTATAACA  AAGGAATCTT  GCAAGGCTAC  CAGATCTCAG
52261   CAATTGTCAC  TATGTTCTGT  AAAAATCACT  TCCTAAAATG  TCTGAATTGA  CTGCTTGTCT
52321   CATTTATTTG  TTTCTCGTGT  CATACTGCAA  TGGATATCTG  TCTTGTTAGT  ATAAATATTT
52381   GTGCATTTTG  TTGTTGTTAA  AACAGCTTTT  TTGGCCTGTC  TTCTTCCACC  TATGAGGTAA
52441   TATAAAACTC  ATGTTTAACA  CTTATTTTTG  TAGCAGGACA  AGCTACAGAC  AAAACCCCTC
52501   AGACACTGAG  TTAAAGAAGG  AAGGGCTTTA  TTCAGCTGGG  AGCTTTGGCA  AGACTCACAT
52561   CTCCAAAAAC  CGAGCTCCCT  GAGTGAGCAA  TTCCTGTCCC  TTTTAAGGGC  TTGCAACTCT
52621   AAGGGGGTCT  GTGTGAGAGG  GTCATGATCG  ACTGAGCAAG  TGGGGGTATG  TGACTGGCAG
52681   CTGCATGCAC  CAGTAATCAG  AACAGAACAG  GGATTTTCAC  AGTGTTTTTC  CACACAATGT
52741   CTGGAATCTA  TAGATAACAT  AACCGGTTAG  GTCGGGGGTC  AATCTTTAAC  CAGACCCAGG
52801   GTGCAACACC  AGGCTGTCTG  CCTGTGGATT  TCATTTCTGC  CTTTTAGCTT  TTACTTTTTC
52861   TTTCTTTGGA  GGCAGAAATT  GGGCATAAGA  CAATATGAGG  GGTGGTCGCC  TCACTTATTC
52921   ACCCCCTTTG  AGAATCTCAC  TCATTAGTGG  GAGTTCTCAC  TTTTATTCTC  ACTACCTATG
52981   TCTTCTTGAA  AGACAGATTG  ATAATGATTC  ATATAGTACA  CTTGTGCTGA  AGCATTTTGG
53041   TGAGCTAAGG  TAGTGATGAA  GCTTTTTATC  ATTTGGAGAA  GTACAGGTAG  CAAACAAGGA
53101   AGCAGTAAGC  AGGTTTCTAT  TAATATTATA  ACTCCTATTA  TAAGAGTTTT  AAATCTTCTT
53161   AGCACTCGGA  ACCATTTTTC  AAACATGGCC  CCAGAAACAA  ATCCATACCA  CACCTACATG
53221   GGCACATGTG  CCACTTTTGT  CATATTTCTA  ACTATGTCTT  CAACTACTTG  CCCTTAATCA
53281   TCTATGTGTA  GACAGCAATT  AGTAAGGTTA  AATTTCCTAC  AGACCCCTCC  TTCAGTTGCT
53341   AGCAAGTAGT  CGAGAGCCAA  TCCATTTTGA  TAGATAGCAT  TTTGCATCTG  AGTTTCTTGC
53401   CAGGCCACAG  TAGTCAGGGC  TCTGCTGGTC  TTATTAGTAA  TTATTTCTAA  GACAGCTTGT
53461   AACCGTATGA  TTCAGTTGAG  CATGTAAATG  GGGGTCCCAT  ATCCCCACAA  GCCGTCTTGT
53521   GCCCAAGTAG  CAGGCCCATA  ATATTGTATG  ATTCTCTCAG  GGGGCCATTC  ATTATTTTTC
53581   CAATTTTCTA  TAGCTATGCT  TTTTTTTTTT  TTTTTTTTTT  TTTTTTTTTT  TTTTTGCGG
53641   GAAGCATATA  CAGGGAAGCC  CAGGAGTTTG  CCTGTCTTTA  TGGGCAGTAG  GAAGAAAGAT
53701   GGTTTAGTAG  TGTCAATAAC  ACAACTACCT  GCCCACTGGT  CAGGTAATTT  GGCATAAGCT
53761   GTATGCCCAC  ATATCCAGTA  TAATCCAGTG  GGGGCTGTCC  AGTCCCGGTG  GGACTCTGGG
53821   TGGGTCCACA  CAGTTTGCAA  CTTTGGGAAT  TTACTAAATA  GATTTTTCTT  AGTGTGGTTT
53881   GAACTCCACT  AGGTGGCTGT  TTTTATAGTA  CTATTATACA  GTTTTTGCCC  AAGGCAGCTG
53941   AGTCTTCCCA  CAGGAAGGGT  GAAGTCCTTC  CCCACTTTTG  CTATACAGTA  TTGTCTAATG
54001   ATTGAGGCTT  TTAGGACCCA  GAAGTTATCA  GGGTGAGTCT  TTTGAGCTGG  GAATTTATCA
54061   GGAACTGGGT  CTGTAGGTAC  TAATTCTCGT  GCTTCCCATG  GCCATTGATC  TCCCATTACA
54121   GTTCCTCCAC  ATACATACAT  AACATGAAGT  GACATTGAGA  GACTGGGCTA  CATGCTCAGC
54181   TAATTGCAAA  AACAAATTTC  TTGTTTTTCC  TGGAATTTCT  AGTACTGGCA  CATTCAGTTC
54241   ATCATAAGAA  GGTTTGAAAT  ACTGGCTCAG  GGGAGCATTT  ATAAACTTCT  CCTCAAACCA
54301   CCATATTTAC  TCAAGGATCC  AGTCCAGCCC  CAACTATTTC  TAAGGTTACA  CGATCCCCTT
54361   TTTTCCAGTG  AGAATCAAGG  GGGTTGGTTA  TTACTAGTTC  TAAGGGGTTA  CACTGACCAC
54421   TGGTACAGGA  AGGGCCACTT  TTCCCTTTCT  GAAGGTGGAC  AGGATTCTTT  TTATTTTTTA
54481   ACCAAGTTGC  CTAAATGACA  CAAGACCAGT  ATCTACATTT  ATTTCCACGC  AGTCTTAATT
54541   CATGACAAGC  GTACTTATTT  TCTGCCATAT  AGCCTCTTTC  CTAATGAACA  GAACCACATC
54601   CTATTTCTAA  CTTATTACTA  TTAATGACAG  CACAGGCATC  AAATTTCAAG  GTGACTTGTT
54661   TGGGCATTCC  TTTTTCTTCT  GTTTTGGCTA  ACACTTTACT  CGTATCGTTT  ATGAACCCCC
54721   ACCAGTCCTC  AGTCCTCAAT  CTTATTTCAA  AAACTGTGGT  CGTGGGAGGC  TCAGATGGGT
54781   CATAACACAC  ATCAGGTTGG  TCATTTCTTG  GGCTACCTAC  CTTGTATAGA  ATAGCATTAT
54841   ACAAACAAGT  TATTTTTAGA  GTCTTTGTAC  ACTTATAATA  ACCATAAAAT  AATAAGACTG
54901   TAGCAACTTT  TTGTCCTACC  TCAGTGACTT  GATGTATACA  CTGGGAACAG  CCCTCAGTCT
54961   GAGGAAGGTT  AGTTGAAGTC  TTTACTGTGC  AAGTCCAAAT  TTTAAGGAAA  ATGAGTCCCT
55021   TGATGAGTTT  TCTCATGTTT  CGGCCATGCA  TGGACCAGTC  AGCTTCCGGG  TGTGACTGGA
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55081   GCAGGGCTTG TTGTCTTCTT CAGTCACTTT GCAGGCGTTG GCGAAGCTGC CACGTACAGC
55141   TCACAGTCTA CTGATGTTCA AGGATGGTCT TGGAAGTTGG GCCCACTAGA ATTAACTGAG
55201   TCCAATACCT CTACTCAGTC ACTTTCAACT GGGCTTTCTG ATACCAGGAG CAAGGTGGCA
55261   GGTTTTAGGG TGTTGCAAAT TTCAATGGTT ATGCAGGGAT TTTCACATAG CAAACTTTGG
55321   TACTTGGTTA ATCTAGCATT TGTTAGCCAA TGATGTATTT ATTAAAGTCA CCACAGCATG
55381   GAGGGCCTTT AAGTTTAGGT TTTGTCCAAG AGTTAGCTTA TCTGCCTCTT GTGCTAGCAG
55441   GGCTGTTGCT GCCAAGGCTC TTAAGCATGG AGGCCAACCC TTAGAAACTC CATCTAGTTG
55501   TTTGGAGGCC CAGCCTCGGC CAGGGCCCCA CAGTCTGGGT CAAAACTCCA ACCGCCATTT
55561   TTTCTCTTTC TGACACATAG AGTGTAAAGG GTTTTGTCAG GTCAGGTAGC CCCAGGGCTG
55621   GGGCCGACAT GAGTTTTTCT TTTAACTCAT GAAAAACTCA TTGCTGTTGG TTGTAATAGA
55681   TGTAGTTTAT CCAATCTACA TTTTTATTAA CTGTCACCCA CCAAAATATT GACTCAAATC
55741   CTGCAGCTAT TTGATTTTGG GATTTAAATT GATCTGCTAT TCCCTGTGGG ACTCCAATTG
55801   CATCTAAATA GATGTGAGAG TTGAAAGACA CATAAGGGTC TTCTCTTGCT TTACGATGTC
55861   TTATTTTTCC TCCCTCTGGT TGATGAAATG CTAGGGTGAA AGGGATAGCC AATTGGACTA
55921   AAGTACAAGT GCCGCTCCAG TTATTTGGCA GAGTGCCCAG TAAAGGTCCA CCACAATACC
55981   ACCACACATC CGCTTGGGGA TGAACAAAGG CTGACTGATT GAGAAGCTCC TGAAAATTCT
56041   TAAGCTCACT GCATCCCTTC AGGTCTCCAA GGAATGCTAA GTTTCCTCCC TGTCATGAGA
56101   GACAAGAAGT GAACTTAGTT TTGGGAGATG GAAGCTGGAT GGCCCTCAGG GGTTGACCTG
56161   CAGGGTGCTG GACTTTGGGA TATAGCAGAG AGAGCTTGGC ACGACTTATT ACTCCAGGCT
56221   GTAGAATCCT GGAAAACAGT TACCATGCAG CCCATGCCTG GTCAACAGGA GGACCACCTT
56281   AGTGGAAAGG GGATAATCTG GCCCTCTGGC CTGCCATGTG CACAAGCATA ACAATTGGTT
56341   TTGTTTAATG TGTGGACAGA ATATTTGATC CATTCCAACT GGGCATTTGC ATCTTGGTAT
56401   CCTGCTTAAT TATCAAAGTT TGTTTTAAGT CTTTAACTTC TATGACCCTC TAGTAAAATG
56461   AATGTATGAT TTTAGGAAAT TACAAAAACC GGTTGGGGCA GTCCATCCTT GCTCTTTAGT
56521   GGTCCACACA ACATTCGACC AACTATGGCA TAAAAGCTCT ACATCGGGGG GCAAGACTCC
56581   TCGTTGACAC TGGGGTCTTT ATTGAAATCT CTCTGGAATA AATGGTCTCA GTTTACTAAG
56641   GCTCAGTCTG AGGAGAGTCA GGAGGGACAG AGGTACTTTT CTGAAGTACA GAGATGTCTT
56701   CGACTTGGCA AGTCCCCACA GGGTATAACA AGGCAAGCAT TAAATTCAAT AGTTTGAGGC
56761   AAAATTGACT TGGTTATGTT AATAACTAGA TGGTCAGAAA TAGAGTGAGG GAAGAAGAAA
56821   GAGTAATAGA ATAGATGAAG GAGTTAAATT TTTCTTAGCT TTAGTTTGGT AGGGTTTTCC
56881   CCTGGGACTA TGGCCCATGA CTCTGGAGGG GGTGGCACTT TCTTGACTCG GGTGTGATGA
56941   GTCCATCCCT TTTTCACCGT ATGAACAACA GTCTCGGTGG TTAGCAGCAC AAGGTAGGGT
57001   CCTTCCTAGG CTGGCTCAAG TTTTCCTTCT TTCCACCCTT TGATGAGAAC ATGATCTTCA
57061   GGCTGGTGCT GGTTTACAGA AAATTCTAGG GGTGGTACAT GTGCTAAAAG ACTTTTAGTT
57121   TTGAGGGAAA GGAAAGTGGA AGATAAACCA AGTATATAAC TTTTAAGAAG TTGACCTTTT
57181   GTTTTAAATG TGGGGACATC AGCAGTGGAC TTTATAGTCC TTGGTGCCTT CTTACTGAGA
57241   AATTTCCTTT AGCACCTATT TTTATTAGTT TTTAGACCAA AGAAAGTCAA ATGCCATTTT
57301   ATATTTGACA ACGCTTCTTG TATGTTTATA CCAGATAAGC TAGATTTCAC CTTTATATTG
57361   GTGTGTTATT AATGTTAAAC TTAGTTTTAA TAAAACTCTG TAGACATATT TATTTGATTT
57421   TTAATGTCTG ACCATAAGGT AAGATTTTTA TAGACTTTTC TTTAACCTTT TATAATTTTT
57481   GTTAAAGAAC AGGTTAGTGC TTTAAGAAAA ACCCGTTGTG TTTTTATTTT AATGTTCAGT
57541   TCACAGAAAA ACTGTATGAT ACCCCTTAAC TTTAGCCAAT ATGTTTAGAC ACAGAATTTT
57601   CTTTACAATT AAGGTTTCAA AACTTGCTTA AACCTTCAAA ACAATTTTTG TAACCTTTTA
57661   ATGTAGGTAA AAATCCACAT TCTTATGCAT CCTCATAATC CTTTTACCAA AGGTATATTT
57721   TACTTTCCTT ACATACCTTG CACATAAACT GTTTATTCAA TAGTTTTACA TTTAGAAGGA
57781   GGCCTAATTA CTTTTAAATT ATACAACATT TCTTACATAA ATTTATTTTT CTAACACACA
57841   TTTTTTTCAT GACTTTCACA GACAATTCTT CGACATGCCT CAACTTTCTG ACTTATTGCA
57901   AACATCCCTT TCTTTAAACA ACTAGTTAAT TTATCTCAGG ACAAGGATTT CCATACAAC
57961   ATTCTTTTTT ATATAAATTC TGCCTCCTCT TTATTTCCTT TTTTTTTTTT CCGAGGATGA
58021   TAACCATTCT TTTCCAAAGC GAACTTCTTT TATGTCTGTG GACTAGACTG TCTAAGGCCA
58081   CAAGATTAGA AGTTACTATA ATACATGTTA CACTGTTAAC TTTTAGCAAA CTTTACTTTT
58141   GTTGAAAACC TTGTAAGTTT GGGATTTCAA TTATCCTTTG CTATTAATAA GACCTTATTT
58201   AGTCCAAATT AACTTAGAAT TGGTATAGAT GGCTTTTTTT TTTTTTAAT TACCTGGGAG
58261   GAACCATCTA TCCTCCTGTC CTGAAGGGAG TTCCTCCTAG GTCTGGTCAG AGCTTTGTAT
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58321   GGTAATTAAG ATTTAGATCC CCTGTTAGGA AACCTGCCGG GTTAAGAGAA TTTTCAGTGG
58381   TTAATGTTAA ATCATCTTCT TTTTTCTTTT TTCCTTAGGA TACTTCTGAA CCGGTGAGGT
58441   GTGCTCACAA TGAGGTTTCC TGTAAAAGTT ATTTTTTTAC TTTCTTCTGT TAGCAAAGCA
58501   GTTGCCGCTA CAGATTGAAT GCATTGGGC CATCCGCGGG TTACTGGGTT AAGGATTTTT
58561   GATAGGAAGG CCTTAATGCT TTTGGAATAT GCCCTGACAA CAAAGTGCCA GTTCCTTCCC
58621   GGTGTTCAGC CACTGCGTTG ATCCTCCACG AGGGCCTGCC ACGTGCTGCT CTGGTGAGGC
58681   GTTCCACCGG GGCAATTGCC TACCTGGGAG CGCTCTCCAG ATCTGTGTCG CTCAAACTGG
58741   CTGGAGTTCC CCGTAGGGAT GCTCCACAGG GCAGGCCTAA GTCGCCTAAG GGGCTGCCTT
58801   GACCGTCCGT TAATCACCTC TGTCTCCAAA AACCAGCTCC CTGAGTGAGC AATTCCTGTC
58861   CCTTTTAAGG GCTTACAACT CTAAGGGGGT CTGCATGAGA GGGTCGTGAT TGATTGAGCA
58921   AGCAGCGGGT ACGTGACTGG GGCTGCATGC ATCAGTAATC AGAACAGAAC AGAACAGCAC
58981   AGGGATTTTC ACAATGCTTT TCCATACAAT GTCTGGAATC TATAGATAAC ATAACCTGTT
59041   AGGTCAAAGG TCGATCTTTA ACCAGACCCA GGGTGCGGTG CCGGGCTGTT TGCCTGTGGA
59101   TTTCATTTCT CCCTTTTAAT TTTTACTTTT TCTTTCTTTG GAGGCAGAAA TTGGGCATAA
59161   GACAATATGA GGGGTGGTCT CCTCCCTTAA TTTAAACAAA ATTTTCAAAG TCCTACCCCA
59221   AGTAAATTGG CAAATATTAA TAAAGTTATG GCATAGAAAA TAAAAATGAT TGTAAAAGGC
59281   GTAAAGATAT TTCTGTGGGG AAAACATTTG TTCATTAGTT ATCAGTTAAA ATTCTGTGAA
59341   AAATAACCAC TAGAGACCCT AAAGTACCCA GGGGCTAATA ATAAGAAGGG AGGAACACCC
59401   TCTCACTCCC CACCGTTACC TGCCCAGAAG GGAAGAGGAA GAGGGTGACT CCAGGAGAGC
59461   TGTGGTCTCC CCTCCCCATA TGTCCACATA TACCTGACCT CCCCTCCCCA AAATATATAC
59521   CCAATATCTC TCCCATATAT ACATATTTAT CTGACCTCTC CACATATGTA TACCTAAACT
59581   TTCTCTATAT ATCCACATAT ACCTAACCCT CTCACACACA TATAGCTGAC CTCCAGTGGA
59641   GGAAAATGGG GAAGAGAGAA GAAGTTATCA AAGGATAAAT CTAGGTCATA CTCAGAAATG
59701   TGAAAAACAA AAACCACACA CAGAAAAAAA AAACACACAC AAAAAAGAAA TTGATAAATT
59761   TGTTTGTGTC AAAATTAAGA ATTCCGGTTC AATGAAGGAT CCCATGGATA AAGTTAAGAC
59821   ACTGCTGTAA GGATGGTAGA GAATTAAATG TCTGAATCAG ACGAAAGGAT GAGTAATTAG
59881   AATGCACAAG GCCAAGAAGA ACAAAACAGA AACTCCACAT AAAAAATGTA TGAGGCCGGG
59941   CGCGGTGGCT CATGCCAGTA ATCCCAGCGC TTTGGGAGGC CAGGGCGGGC CGATCAGGAG
60001   TTTGAGACCA GGCTGGCCAA CATTGTGAAA CCCCATCTCT ACAAAAAATA CAAAAAATTA
60061   GCCGGGCGTG GTGGTGGGTG CCTATAATCC CAGCTACTTG GGAGGCTGAG GCAGGAGAAT
60121   CACTTAAACT CAGGAGGCAG AGGTTGCAGT GAGCTGAGAT CACACCATTG CACTCCAGCC
60181   TGGGTGACAG TGTGAGACTC TGTCTCAAAA AAAAAAAAA TTATATATAT ATATATATAT
60241   ATATATATAT ATATATATAT ATATGAAATA AATGAACAAG AAATTTAGAT ACAGGAAAAT
60301   CCAAAGCACT TGGTAATGAA AGAAAGGTAA AGTGATGTGT CCTTTTGCAT TTAAAAGAGA
60361   GCATTAACAA ATTAGAGAGC TGAATAATGC TCAGTATTGG TGTGGATATG GAGACTCAGG
60421   AATCCTCATA CACTGCTGAT GGGAGTGCCC ACTCCCTGGG AATATTTTCC AAATATCATC
60481   TCAAACATAT CCCATAAAGG TGACAGGAAA GTGTGGGCTG ACTGATATCC TTCACTGAGA
60541   GAGGTGGAGG TAAAATGAAG TCACTGCACA ATATAGAGTT GGAAGCAATG GATTAGATGT
60601   CCACATAGTT ACGTGGAAGA ATCCGTAAGA TACACACACA CACACACACA CACACACACC
60661   TTTGTGTATA TTGTTCCTGG CAGGTAGGCA TGGAGGTTTA GAGGCTTTCT ACATCACACC
60721   TACTGCACAC AGTAAATGGC CAGGCTGAGC ACTGACTTCC ATGAAGGGAG ATTGAAGGTA
60781   AGAGATTGAA GATTGTTCCC TGGTCTGGGA CCCTGCAACT GAATATGCAG AAAAAAGTAC
60841   ACCCCGCCAC CCCGCTTCCC ATCTTTCCTA CCTGATTAGA ATAGCTTTTT CAGAAAACGT
60901   TGGCCAGGGG TTGTGGCTCA CACCTGTAAT CCCAGCACTT TGGGAGGCTG AGGCGGGCAG
60961   ATCATCTGAG GTCAGAAGTT CCAGACCAGC CTGGCCAACA TGGCGAAACC CCATCTCTAC
61021   TAAAAATATA AAAAATTAGC AGGGCATGGT GGCACACACC TGTCATCCCA GCTACTCGGG
61081   AGCCTGAGGC AGGAGACTCA CTTGAAGCAC AGTGATGGAG GTTGAAGTTA GCTGAGATCT
61141   TGCCACTGCA CTCCAGCCTG GCAACAGAG TGACACTTTG TCTCAACAAC AACAACAAAA
61201   CCCACCAAAA CTTTAAATCT ACCTATGGCC AAATGCCTGC TAAAATGAGC ACCCAAGAAG
61261   CAGTGTTCAG GAAAGTCAGA TGAATACCCT AAAATTAGAT GCAATGTTGG CTGGTCACAG
61321   TGGCTCAGGC CCTGTAATCC CAATCCTTCT TGGGAGGCCG AGGCGACAGA TCGCTTAAGC
61381   TCAGGAGATC GAGACCAGTC TGGACAACAT GGTGAGACCG TGTCTCTACA AAAACGTACA
61441   AAAATGAGCT GGGAGTGGTG GCGCGCACCT GTAGTCCCAG CTACTCAGGA AGCTGAGGTG
61501   GGAGGATCTC TTGAACCCAG AAGGCGGAGA CTGCAGTGAG CAGAGATCAT GCCACTACAC
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61561   CCCAGCCTGG ATGATAGAGC CAGACCCCCA TCTCCAGAAA AAAAAAATAA AGAGAGAGAG
61621   AGATGCAATA TTTAGGGTTC AACAAGACTG AATTTCTGAC TCCTTTCCCT ACCTCTCCAG
61681   CATGTTAGAT TCTGGGTCCT TCATCCTAAC CCCCTGTTCA TGCCATAGCC ACCCTGTGGT
61741   ACCAACTTTG GAAGCCTGGA TCTTCATCCC CTCATGATAA TGAGTGTCCC ATCAGGTCTC
61801   CATGCTCAGC TTGGCAAGAG TATCTGTCTT CTCCTCATGG GACGGTCACA TTCACCCAGC
61861   ACTGACAGGT TCCATTCCCA CTAGGGTGGC ACCCTATATG GTCTGAGTCC AGGCCTTCCT
61921   GGTCCCTCAG TAATCTCAGC ATGGTAGCAC AATCGAAAAG GGCTAGGCAC GGCAGCACCA
61981   TTTCCCACCA AGAGGTCTGA TGGCTCATCA CATAGACTGA AGGAGATTCT GAAGAGCAGA
62041   GGTGGAATGA AGAATGAATC GTGGGCTCTG CTCTTCCTAG GCCTGTCTTC CTCTCTCCCG
62101   AGATGTTAGC TAACTCATGA GAGCCAGAAA CCAACTGCAG GCTGGCCTCA GGCACTTAGG
62161   TAGTGCTTCA GCCTCAGCAG TCCACATTCT AGGAACCCTC ATAATATGGG TTGAAGTATG
62221   CATTCCCACA AAAATAAAGT TGTTGAAGTC CTAACCACCA GTACTGAAAT GGGAAAAGTT
62281   CCCTTGTCCC GCTCGCATGG CATGTGATAG GAGTGTGGCT AATTTCTTCA GTGCCTGGCT
62341   GCTCAAACCT CTAGGGGAAC ATTAAGACGG GCAGGTTGTG GGTCTCCAAC CCCATGACCC
62401   CACCACAGTG TCTAGGGTTG AATGTTTACA GCTCCTGAAG CCACAGTGGG TGTGTGTTAC
62461   AGGGTGCTCT TTTAGTTTTG CCATTTATAG GCAGCTGGTG TTAACCAACT CAATTAGACC
62521   GTCTACCTTG TCCCAAGGAC AGAAGAAGGC TTTCTGTATC CCAGGTTCTT GCCTTGGTGT
62581   ACCGGAATAA ATCAGACCAC ACCTGGGCTT AGAGAAAGAG TGCAAGGTTT TATTAAGTGG
62641   AGGTAGCTCT CAGCAGTTGG GCAAAGCCAA AAGTGGATGG AGTGGGAAAG TTTTCCCTTG
62701   GAGTCAGCCA CTCAGTGGCC CAGGCTCTCC TCCAACCACC CCAGTCAAAT TCCGCCTCAT
62761   TTTGCCAGGC AAACGTTTGT TGTGTGCTCT TCTGCCAGTG TGCTCCCCTG GACGTCCAGC
62821   TATTCGTGTC TTGTGGCAGG CCAGGGGAGG TCTTGGGAAA TGCAACATTT GGGCAGGAAA
62881   ACAAAAATGC CTGTCCTCAC CGTGGTCCCT GGGCACAGGC CTGGGGGTGG AGCCCTAGCC
62941   GGGGACCACG CCCTTCCCTT CCCCACTTCC ATATCATTTA AAGGGACCAT GCCCTTCCCT
63001   TCCCAGCACT TTCCCCCTCC TGTATCAGGA CCTGTGAATG TGGCCTTATT TGGAAATAGG
63061   GTCTTTGCAC TTCATCAGTT AAGATAAGAG TGGGCTCTAA CCCAACATAA AGGGTGTCCT
63121   TATAAAAAGG AGAAATGTCA TACACAGAGA CTGACACCTA TAGAGAGAAA ATGTGGTGAG
63181   TAGACACAGG GAGAATCACC ATTCAAGTCA AGCAATGAGT CTGGGGATAC CAGAAGCTGG
63241   GAGAGAAACC TGGAACAGAT TATCCCTCAT TGCCTTCAGA AGGAATCAAA CCTGATGATA
63301   CTTTGATTTC AGACTTCCAG CTTCCAGGAC TGTGTGACGA TAAATATCTG TTGTTAAGCC
63361   AACGAGTTTG AGGTACTTTG TTACTGCAGC CCCAGAAAAC TAATACAGTA GGTACTATGG
63421   ACTGAATTGA CTCCCCGTCG CAAAATTCAT ATGTTGAAAC CCTAACCCCC AGTGTGATGG
63481   TACTTGGAGC TGGGGCGTTT GGGAAGTCAT TATATTTAGA CAAACTCATC AGGATGTGTC
63541   TCTCATGATG AAATTCATGC CCTTATTAAA AGAGACAACA GGCCAGGTGC AGTGGCTCAT
63601   GCCTGTAATC CCAGCACTTT GGGAGGCTGA GGTGGATGGA TCACCTGAGG TTGGGAGTTT
63661   GAGACCAGCC TGGCCAACAT GGTAAAACCC CATGTCTACT AAAAATACAA AAATTGGCCA
63721   GGTGTGGTGG TGCACGCTTG TACTCCCAGC TACCTGGGAG GCTGAGGCAG GAGAATCCCT
63781   TGAAACCAGG AGGTGGAAGT TGCAGTGAGA TCACACCACT GTACTCTAGC CTGGGTGATA
63841   GAGACTCCAT CTCAAAAAAA AAAAAAAAA AGACAATAGA GCCAGGTGCT GCAGCTGATG
63901   CCTGTAATTC CAACACTATG AGAGGCTGAA GCAGGAGGCT CGCTTTAGCC CAGGAGTTCA
63961   AGACCAGCTT GGACAAAATA GTGAGACCCC CAACTTCTAA AAATTTAAAA AATGAACTGG
64021   GTGTGGTGGT ACACATCTGA GGCTCCAGCT ACTCTGGAGG CTGAGGTGGG AGGATTGCTT
64081   GAGCCCAGGA GGAGGCTGCA GTGAGCCATT GCTGTCCAGC CTGGGCTACA CGAGAACCTG
64141   TCTCGGGAAA AGGAGAAAAC AGTGAGACCT CTTTTTCTCT CCTCCTTCTC TCCACTGCCT
64201   AAGCCCTACA AGCACAAAAA GGACACCACA TGAGCACATA GTGAGAATGC TGCTGCCACC
64261   AACAAGTCAG GAAGAGAGCG TTCACCTAGA AACTGAATTG GCCAGCACCT GGATCTTGGA
64321   CTTCTGAGCT TCCAGAACTG TGAGAAAGTT ATTTTTTTTT TAGCGACTAA GTCTATAGTA
64381   TTTTATTACA GCAGCTCAAG GTAACTAACA TAGTAGAAGG GATGAATTAT GGAGATCACA
64441   AGTCCACGCC TCCAGAAAAA GACTTCCCTA AAAATTAGTC TGAGCAAAAT TCGAATGATG
64501   AATTATTTTT AAGAACTTTT AAGGGATCTG ACAAGTTTGC AAGAGCTAGA GAATGCTTTA
64561   CAACGTGATA ATAGAATGCT CTGTGATGAC AGAAATCTTT CCACACTGTT CAAAACTAGC
64621   TACTGGCCAC TTGTGACTAT TGTGCACTTG AAATGTGACT GGTGTCTGAG GAGCAGAATG
64681   TTTAATTTTA CTTAATTTTA ATTCATTACA ATAGCTACAT GTAGCTAGGG CTACTGGAT
64741   TGAACAGCAC AGCTCGAGTC TTTTAGAGGG AGACAGGACT CACCAAGATG GATGCTGGTG
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64801    GCCAAGCAGC AATGGCAGGT AGTACACACA CAAGAGGCAG ATGATACAAC ACATCCTTCC
64861    CAAACCTGGA GATAAGCTCA CCCCACAATC CCGCCGCTGA AATAGAGTTG ATGTTACCAA
64921    TGTGCATTTT TATGTCCTTT TCCATACAGA AAGATCATTC AGCAAGTACT ATGGTACTTA
64981    AAAAACAACA TTCAATTCAT TATTATGACA AAATTAAATT AATAGCTCTT CCTTAAACTT
65041    TTAAATTCAA TTTACAATGC TTACTATTGG CATTTATTAA TCTACCAATT TTTTCCCATA
65101    GAACCCATAG AACAAATAAT CTACCAAATT TTTAACATTC ATTTTTGGCA AGGCTTTTGC
65161    AATTTGACGA ACTTTAAGAA GAAAACTTAT AAATTGCAAT TTTTAAATCT GACATACTGG
65221    ACTTTTAAAG TATCCAATTG ACTAATGAAC AAAACTGCTC CAAATTTTTC AATTCTTAAA
65281    AATCTTAAGA CAATACTTAA TATGGCAAAT CTTAACTTCT TAAACTTTGT AAGAATGCTA
65341    ATCAACTTAG ATTGGTATAA AGTTGAGTTA AAAATCACAG GATACATCAT CTCAGCTATA
65401    AGTTTTCATG AGTTGAGTTT TTACAATCAC TTGAAATGCT TAGAATAGGA AATACGTATA
65461    AATTATTTAA CATAAAATAT TGTTACAAAA CCTCTGGAGT GTCAGTTTCT CTGGCCAGAC
65521    TTTATGCTGC AGCACCTTTG CCTGAGTTCT TGTCCTGCAT CCAGGAAGAA TTAGGTACAG
65581    AGGCAAGAGT CAAGAAGATT AGTTTTCCAA TAGTTCAGCT CACCTAGTTA ACTCCTGTTC
65641    ACAATCTTCA AAGTTATCAG AAACCTGCAA TTGAGGGTTA TAATCCATTC TTTGCAGAGT
65701    TTCAAAACAA GACAACATTT GTCATGAAT GTTAAAATGT CCTAGGGTAG TCACAGTCAA
65761    AAACACAATT GACAAAGAAA TTTAGTCACC TCTGTGATTT ACAATAGCCT AACACAATAA
65821    CTCTAATTAT AACTGATGAC ACAAACTCAG ATATCAGAAC TCTAGAAATC CCCTATAATT
65881    TTGGAACACA CATTCACAGT TTTCACTGAA ATATGACCTG AAGATCAAAT ATCACCTTAT
65941    TTCAACAATC CTATATAACT AAACGTGTCA AATGATCCTG TTTACCTCTC CTTTGGATAC
66001    TCCAGGGGCC CTCTGTAGCA TCCAAAAGTT AGGGGTTAGC AAAGACAATT TTGAAGCTGT
66061    AAAGGCTCAA AACACTTAAT GAACCTCTAG TCATATCTGT TCTCTACTCA CTAAATGCTA
66121    GTAGCACCTC TCAGTTGTGG CTAAGCTGGG AGGATCTCTT GAGCCTAGAA GTTTGGGGAC
66181    GCAGTGAGCT ATGATTATGC CACTGCACTC CAGCCTGGGC AACAATGCAA AATCCTGTCT
66241    CAAAAACAAA AACAAAAAAC AAATTGCCTA TGCTGTGGTT ATCTCACAAT TAATAAAAAG
66301    GAAAAAAAAA GTATGCAGTC TTTGTAGGTC CTTGGGGTTT GTTGGAACTC AGAAAACAAT
66361    ACCCCAAAAT AAAGACCGCA GAAGCCAAAG TTTTTCTCTG ATCTTCTCCT GCCCTCCTGT
66421    CTCTGAGTCC CATTCTCCCC GGAGTCTAGC CATAGAAATG AGAATTCCTC TTCCTCAAGT
66481    TAGGTCATAG AAATCAAAAC ACCTTTTCCC CAGAGCCCAG CCATAAAACC TAAAAATATT
66541    ACTCTAACTT TCCCTCTGTT TTTCTGTGTA AAAACTGGCC ATAAAGAAAT TATCTGAACT
66601    ACCTTATTTG ATCATAGATC ACCAGACCGC ATTCCAGAGA GGATCCAGAA GGAAGGAATG
66661    CTGCACAGAG AGGCGAAGAA GAATCTAGAC AGACAGGCCT TGCTGGGTTT CCCTACTCTG
66721    TTTATTAGCA ATCCTATTTC TACACGGCGG CCCATACTTT GTTGAATCTA AAAAATAAAA
66781    ATGGACAATT TCCCCTGTAC ATGTTAATAC ACATTAATAA ATTGGATATA AATTGGATAA
66841    TTTATTAATA TACACATTAA TAAATTGGAT GCAGCCGGGT GCAATGGCTC ACGCCTGTAA
66901    TCCCAGCACT TTGGGAGCTG AGGCGGGCAG ACCACGAGGT CAAGACCACC CTAGCCGAAA
66961    TGGTGAAACC CCGTCTCTAT TAAAAATACA AAAGTTAGCT GGGCGTGGTG GCACATGCCT
67021    GTAGTCCCAG CTACTGGGGA GGCTGAGGCA GGAGAATTGC TTGAACTCGG GAGGCGGAGG
67081    TTGCAGTGAG CCGAGATTGC GCCACTGCAC TCCAGCCTGG TGACAGAGTG AGACTCCGTC
67141    TAAAAATAAT AATAATAATA ATAATAATAA TAATAATAAT AATAAATTGG ATGCATTTTA
67201    TCCTATTAAT CTTCCTCTTG TCGGTGGTTT TCAGCGACTC TTCAGAGGCC AAAGAGTAAG
67261    TTTTCCCTTA GCCCCTACAG GTTCTTATGT TTAATTTGTT ACTCTCATTT AAGACATAAT
67321    TAAAGTGGCT TCTCCATGAA GATTATTTCT GCATCCATTA TTTGGTAAGA TTGGCCGTTT
67381    TCTCCTTTGA TCTCTACTTC ACACTGACCC ACATAAAACA TCACTGCCTG TTTTTTTGTT
67441    GTTGTTGTTT GGAGACGGAG TCTTGCTCTG TTGCCCAGGC TGGAGTGCAG TGGTGTGATC
67501    TCCGCTCACT GCAAGCTCCG CCTCCCGGAT TCACGCCATT CTCCTGCCTC AGCCTCCTGA
67561    GCAGCTGGGA CTACAGGCAC CCACCACCAA GCCCGGCTAA TTTTTGTATT TTTAGTAGAT
67621    ACGGGGTTTC ACTTTGTTAA CCAGGATGGT CTCGATCTCC TGACCTCGTG ATCGGCCCGC
67681    CTCAGCCTCC CAAAGTGCTG GGATTACAGG AGTGAGCCAC TGCGCCCGGC CCCGTTTTTT
67741    TTTTGGTTT TTGCATGTCT TCTCCCTTTT ACTGTAAACT ATTTCCACTA CCAGCGTAGT
67801    TATCATTTCT ACTGCTTAAT AATTGTTTTG GGGAAGTGAA TGCATCAACC CACATGAATT
67861    TCTTGTCTAT TTGACAATTT ATTCTCTTTA GGAATAGTAT TAACTCCTAA GGTCCTGGGA
67921    GCCAGTCTCT GTACTTGGCT GCTCCAGGGT CCTACTTCAG TTTCCCAGCT TCTCAGTACT
67981    GTCACTGTCA ATTGTGGGTA ATAATTATTT TTGTCCACCA AAAGACTCTG TATGTGAATG
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68041   AGTTTTGAAA TCTGCTGAGT AATACAGTGT CAACCCAGTT AATGATTTGC CGGGCGGCTT
68101   GATCAGGGGC TGTCCAACTA CCGGCATTTT GATTTGGAGC GTCATCTAGT GTCTGAAAGC
68161   ACAAACAACA TCCTACATTG TAAATGCCTT TGGCTACAGA GATTGAAACC AAAGCAAACC
68221   TATGTTTTGA ATTGTTATTC TTCAGCAGTT CTGCTAGCTT TGAAAAATCT AAAAGTTAAA
68281   AAAAAGCTTT ATATTTCATT TTCTGCCTAA ACTCTTTAAA ATTGCTAGTT GACAATTAGA
68341   TATTTTCAAT TTAATGAAAT TTTTTTTTAG TTCACAGATT AATACACAAT GGGGGAGGGT
68401   TCTTATTCTG TTGGACTTTT ACATAACCTC CACTTTAGTG CAGTCTGCTT TATGGGGTCT
68461   TGTTTGAGGT GTGTGTGTGT TTAAGGGAAT GTGGTTTACA ATCAAAATAT TGGGTTGCTC
68521   TTAGGCACAT TGTAAAGTCA CACACCTGTA TTCTTATTGA TACATAATGA TTAATAACAT
68581   TATTATTACA GCCTGATCAC CATCATTATT GATATATCTA AATAATGAAT TTTATAATTT
68641   TGCTTCCTGT CAGGCAAGAG CCAATTTCAG TGCTACCATG TTTGTATAGC AGTATTTATG
68701   TCTGTCATCC TCAGTCATTT TACTTCACTT GTTCTTAGCC AAACGGCCGA GAAGCGATGG
68761   TCATTTTACT TCAAAAATGA AAAGAATTAA TATTTTTACG TTTCCCTTAA AGACCCTATG
68821   TTTAACCTCC ACTCCTGGGT AAAATGGTCT AGTCCCTCCT TTTCATATCA TCTCTGATAT
68881   CTTTTGCACA GCCACTATTA CCTACCGTTT TCTAGATCCC TATTCTTCAA ACACCACCAT
68941   GAAGGTAGAG CCTGTCTGAA TTATTTTCTT GTCCCCTGAA CTCAGTACAT TGTTAGGCTT
69001   CTTGAAGATG TTGATCAGTT GTTTGTGGAG TGAATGAATC AGCTAGCATG ATTTTTCTAG
69061   ACCACTGAGA CAAGTGTCTA AGACACTTGT TCCTTCCCAT GTTCTTGCCT GCCTGTGCAA
69121   TCCATGCAGT CTCATGGCTT CCCAGTGCCT CAGAATTATC CCCTGTCAAA CAGGCATTAT
69181   AATTTCTGTC CACTGAAAAG GACAAAAAC TAAGTGTATA GCTAGAAGTT AAAAATTACC
69241   GGCCAGGTAC TGTGGCTCAC TCCTGTTATT CCAACATTTT GGGAGGCTGA GGCGGGCAGA
69301   TCACCTGAGG TCAGGAATTC GATACCAGGC TGGCTAACAT GGCGACCCCG TCTCTATCAA
69361   AAATGTAAAA GTTAGCCAGG TGTGGTGGCT CGCACCTGTG GCCCCAGCTA CTCAGGAGGC
69421   TGAGGCAGGA GGATCGTTTG AGCCCTGGAG GTTGAGGCTG CAGAAAAATA GGAATATACT
69481   CTCTTTCAAG AGTTCGTGGT TTTGACTGCC ACCTAGCGTA CATCAGAAAA ACCGCATGAC
69541   ATAGGAAATG CCTGTGACAG AGGGGTAAGG TGAGAGAGGT TGATGAAGAA TGTATTGAAG
69601   GAGTGAAAAC GCTTCCATCC CTCTACTTAC TAAATATATT AGTTAAGTAG TTGGGGCATA
69661   TTTTAATTCA TGCATTTTGT AGATAGAAAA ACAAAGTTT TATTCTGTTT GATTTAGTTG
69721   ATACTTTAAT ATGTGTGTGT TTAGGATGCA TGATTTATAA TCAGTCTGCA GCACTTCTTG
69781   GAGAAGTCTG AATTCTCATT CTCCATTTCC TTATTGGCAA CGTGAGAATG ATTACAATGG
69841   TGGTTGTCTC ATAGAATGCA GGGAGTCAGA ATGAAAATAG TCCATATAAT GCCTGGTGCA
69901   GAGGAAGGGT TCAGTTAACT GTCTGTATTA ATATTACTGA TAACAGTCAT GACAAACAAA
69961   AGCTTAACAA CAACACCACC AACAACAGTT GCAGAATTGA GCCACCAATT TGCACACAAG
70021   ATTGTAGGTA GGATGTTTTA GAAAAGTTAT TATTTAATAT ATGTATATAT TTTTGTACTT
70081   AAAATATGTC AGAGGTTGTT CTAAGAACTA TTTAAATGTT AACTCCTTAA TCCTCATAAT
70141   GACCCATGAA ACAGGTAGGC TTATTATTGT CTCTTTACAT GTGAGAACAC TGAGACACGA
70201   AAAGGTTTAT TAACTCACCC AAAGTCACAC AGCTGGTAAA ACGGCAAAAT TGAATTTGAA
70261   CTCAGACATT CCAGGTTCCA AGACAGTCTA ATTATTCTTT TGACTAATAT ACTAAGCTGC
70321   CTCTGTATTT TTCCTTGATT ACTTTGTAAA AGTATGAGGA AAATATAAGT GCTTCAAGTA
70381   ACCATGAAAA ATATAAACAA TCTATGTATC AACTGAAGCA TAATTACAAA TCCTTTGATA
70441   AGCAAACATA ATAAAAATTT GATATCAATC AAAACTTTCA TGTAATGTAA GCAGGTTGAG
70501   ATGAATTCTA TAGTAAAAAA GTGCAGAGTG CTGGAATACC ATGCTCCTAA TATATTGGCT
70561   AGGCACACCT GCCTGCTATC AAAGGTATGC ACACACCTTG GATACAGAAA GTTGGGACTG
70621   GGTAGTTATG TGAGTGTCAT CAGAATTCTT TCCCACTTGG GAAAGAATTG TCCATCATAA
70681   GCTTGGATGA TGGACAAGGA GTGAGCTCCC AGAACAGTGA TGTGGGGATA CATCCTCACA
70741   TCACAGTGAG AATGAGTGTT CTAGACTGTT TACACACCTA CCACTCCTAA ATGCACACAT
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70861   TCTTATCATT AGGCTTCTTG GGCTAGTAC CTAGGGCCTG TATCCTTTCA GAGGCAGCTA
70921   AGGGAAGCAC ACATAATTAG AAAGAATGAA CCAGCTTGTT GGATTTGGTC TCTTCGCATC
70981   CAGCCCTCCA AGTTAAGGAG AGTACCATCT TTCTTAGGGT CACCAAAGGA AAAAAAAAAA
71041   AAAGAAAGAA ACAGAAGGAT ATCATACAGC AAGGATCTAA TGCAAATATG CCTCAAATGA
71101   GAGGCTACTG TGTGCTGATC CCAATCCCAG GAACTGTATG CACATTATCT AATTTAATCC
71161   TCACTGTATT TCTGGGAGTA TTATTCCCAT TTTACAGAGA AGGAACTTGG CAGGGTAACC
71221   AAGCTCATGA ATGGAGAAAC TGGGATTAAA TATAAAGCTT CCTTGCTCCA GAACTGCTGT
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71281  CTTTCTGCTC TTCCACACTA CCAGCTCAGC TGTGCTCTCT ACATGCAGGC AGTTTTACAA
71341  GTTTCAGATT AGCCTGGGAC TTCCAGGGTT TTGAATGGGT TAGGGAATGG GGAACTTTTG
71401  GGTTTACTTT CCATTTTTTC TTCATACATA TGTAATATAT AACATAAATC TATGGTATAT
71461  ATGATAAATA TATGGCTACA TATGAACTAT ATAATCACAT ATATGCATTA TAAATAAATA
71521  TTAATTTTAT AATATTTTAA AGGTTATCAA ATAAATATTA ATATAAATAA TTAAATAATT
71581  AATACTCAGC TTTGTTTTCC AAAGTGATAA ATGCCTATAT TTAGCAAAAT ATTTTTTGGA
71641  GGCCTGATAG TTTTTAGGAG TGTAAAGAAG TCCTGATATC TAAATGTTTA AGAACCACTA
71701  TTTTAGGCTG TTGTCTTCTG TCTTATTTTC CCAGCTAGAC TGGTAAATAC TTGAAGGCAA
71761  ACGTTTAGCC AGCACATTAA CATTTTATGT TTTTATTCTT TTGTGCTCTC AGTGGCTGTG
71821  TCTTTTCTAT CGATTTCTCA CACTGTATGA TGGTTATATT TGTCTGTATC TGTCCCACCA
71881  GGTATAAGTT CTTGAGAGGA CACACTGCTA GGCTGATCTT AGTTTTTATT ATTTCTCCTG
71941  GTGTCCTGTG CTTAACAAGT GCTCATTAAG TGTGTAAAAA CACAGCACAG TAAAAAACTA
72001  GACATTAAAA AATAATGTCA ACCAATCTAT TGAAATTTGC ATTTCCATGT TTCTTCCAAT
72061  ATAGTCATTG TGTCAGGTTA TGTACTTATT CTGATGAAGA CTATTGCCTA ATATACGTTT
72121  GCATCTTGTG CTTTATAACT GCCTTCATAT AGACACAGAT TGAGAAGGTG TAAAAATGTG
72181  CATATCCTCA CAATTGACAA ATTCTTATCC TTTGAGGGTA GGTTTGACTT TCTGAAATGC
72241  TTTGACATCA TTTGAAAGAA GCTTGAAGAA TAAGATAGCT GTTAATGACC CAGTTTCCTA
72301  TGTCACTTAT ACAATTATAA TGGCAATTTC AAAATGTTAG GTAAATATAT TTTGCAATAT
72361  ATTGTTCCTT TTGTAATACT CTCTATGTAT TTATTTATAT TTTTAAATTT TATATTTATG
72421  TATTTATTTT CTGGACAGA GTCTTGCTCT GTTGCCCAGG TTAGAGTGAA GTGTTGTGAT
72481  CATAGCTCTC TGCAACTTCA AACTGCTTGG CAAAAGTGAT CCTCCTGCCT CAGCCTCATG
72541  AGTAGAGTAG CGGGAACTAC AGGCGCATGC CACTGCACCC AGCTAATCAC TATTTATTAT
72601  GCTCCTACTG TGTGCTTTAG TATATTTTCT GTTGTTTTCT GCAACCCATT TTGAGGGCGT
72661  GTTAGGGAAT ACAGATGCAG TAACTTTCGT CTCAGCCCTT GAGGTGAGGA AATATTTAGC
72721  CTCAGGTTTA ATCTAATTGT TGGCCATTTG CCTTCAAAGA TTGAAATATG AGCAAAACTG
72781  TGGCTCTGGG TTATATGTTA AAAAAAAGTT TATGGGGCTG AAGCCAGGCA ACAGACAAGA
72841  GCCCCTACAA TCTTATTTAG GCTGAAAATA TCCTGGAGTC CCTGTATTGT TGGTCTCAAG
72901  CAGATAGCAA CACTAACACT TACTCTTTGA GGCAGGCACT GCCAGTGGGG TGGCTGTTAT
72961  TATTAGCTTC ATTAATTGGT GAGTCAGGAA AAAACAGCTT TAAATCATTC AAAGTTCTGG
73021  CCTATACAGG ATTTAGTAAT ATTAGGTTAG CTACATCCAA AAGATGACAG AACCCTACTC
73081  TAAGGCTGGG CTTGGTGGTT CACACCTATA ATCTCAAAAC TTTGGGAGGC TGAGGCAGGA
73141  GGATCACTTG GTGCCAAGAG TTTGAGACCA GCCTGAGCAA CATAGTGAGA CCCCTGTCTC
73201  TATCAAAAAC AAAGAACTCT AATTGGCATA GTAGAAGGAA AAAGTGAAAG AAAAACCAGC
73261  TGTCACCCTC ATTCCTTACA CCTGTCCTAA CAACTCCTCT CACTATCCTT TGAATATATC
73321  TTGGCTGTTT GAGTCTCTCT CTAGCCCCAT TACTGCTGTT TGGACTTGAC ATTTTGCTCT
73381  GCATTTTTAA CTTTTCTACC AGGGTTTCCA GACCCTGAAG AGTGTGGCAT GAAACAAAAC
73441  TAGTCAACCT ATAATATTTA TGATGTGTGT GTAAATAAAA GAATACACAA TATATTGCAT
73501  TACAATATTT TAACTGTGTC CTCAATTTGT TTGTGGCTTT CTTGAGGACA TCAGTTTTGG
73561  GTGGGACGAC CACATCCTTA ATCTGAACTT TCCCTTGGAG GTCATTCTTT TTTTTTTGAA
73621  ATAGAGTCTC GCTCTGTCAC CCAGGCTGGA GTGCAGTGGC GCAATCTCAG CTCACTGCAA
73681  CGTCCGCCTC CTGGGTTCAA GTGATTCTCC TGCCTCAGCC TTCCAAGTAG CTGGGATTAC
73741  AGATGCACGC CACCATGCCG AGCTAATTTT TGTATTTTTA GAAGAGACGG AATTTCACCA
73801  TGTTGGTCAG GCTGGTCTTA AACTCCTGAC CTCATGATCT GCCCACCTCA GCCTCCTAAA
73861  GTGCTGGGAT TACAGGCGTG AGCCACCCCG CCCGGCCAGA GGTCATTCTA ATAGACTTTT
73921  TTTTTGTTGT TGCTCACAGG CTTGTTCAAT CTTATTTCAA AATTTGAGAA ATACAGTTTC
73981  CATGGAACAC CAACCAGATA TCAGGTTGCT ATGGAGTTGA TAGTCAAAAG CTTTGTATCT
74041  TCCAGTTTTT CAGAATGGCT TCTAAAGGTT CTGATTCAGA GCTCTTAGGC GAAATTGAAC
74101  AACCAAGTGT CAAAGTACAA CATTCAGGAA GTTAAAAACA TGACTGACAT ATATGTACTA
74161  TATATAGTGA GCTTGTGTAT GTGTCAATGA ATGATTTAAT TCATTAATGA AGGAGGAAGC
74221  AGAATCACAA TTAGGTCAAA GGAAGATACG GGAGAATAAA ATATGTATTT GGTCAGGGAA
74281  AGGATGTATA CTGGAAGAGG AAGGGAAAAT CAGATATAAA GTTGTTTAAT GACTTATTAG
74341  GCAATACAAT AATAACTTTT AGGGTCATTT TTTCTATATT AAGAATTCAT TTCCATCTCT
74401  ATGACAAAAT CCTTATTAAT TTATTAAACT TCTACAAGTG AATGTTTACT TTTAGATAGT
74461  CTGGACCCAA TAAAATGTAA ACATTAAGTC AGAGTTACTT TCACGTAGGA CAGTGTTGTC
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74521   CAATAAGGTA CCACTAGCTA CACGTGATCA TTGACCATTT GGACTATAGC TAGACTGATT
74581   TAAAATGTTC TAAAAGTGTA AAATACACAC CAGGTTCTGA AGATTTATCA TTTAAAAAAG
74641   AATGTCAACT GTCTTTTTTT TTAGCTTATT TATTATATGT TGAAGTGATA ATAGTTTAGA
74701   TATATTAAGT TAAATAAAAT ATCTTAAAAT TAATTTTACT TGTTTCTTTT CATTCTTTCA
74761   ATGTGACCAC TAGAAATCTG GAAAGTATTT ATGTGATTCA CATTCTATTT TACTGTCTAG
74821   TATTGCCTTA CATCATCAGG TACCCCATAA GTAGGCTTTT TAGATAATTC TCTAATATAG
74881   CTTGGAAGGA TATGGAGAAA TATTTTTGCG TTGCTTTTAA GTTTTGCATA ACTTTTTCAA
74941   CACACTTTAT AAAGGATCTA GAAAAGGGTT GGTTACATGT TTCTCTGTCT TCTGGCCTCC
75001   ACCATGTTGC CAGGAGGTTG GGGACAAGAT TCTGGGTGGC TGGATGTCCT AATGGCTTGA
75061   GGTCTGGACT TGAGATTTGC ATATAAAGAG ATGTGATTAG ATTGAGTCGA CTAGAAAAAT
75121   CATATTAGAG AACTGAATCA CAGCGATTAA ATTTACATGT CGATTTATAA ACCAGGACAC
75181   CAATTTATAG TGAAAGAAGG TCCAGTTACC TGGTAATCAA GACGTTTCAT AGCTATTTTC
75241   ATGATGGATA TACTTAGCTG AGTTTTAAAT GAGAAGGGGG TTCATTGCAC ATAGAATAAG
75301   ATCTAAGTGA AATGTTTATT TTATTTTTTT TTTTTGACA TGGAGTCTTG CTCTGTTGCC
75361   CAGGCTGGAG TGCAATGAGG CAATCTCGGC TTCTGGAGTG CAATGAGGCA ATCTCGGCTT
75421   CTGGAGTGCA ACGAGGCAAT CTCGGCTCAC TGCAACCTCC ACCTCCCGGG TTCAAATGAT
75481   TCTCCTGCCT CAGTTTCCTG AGTAGCTGGG ATTAGAGTTG CCTGCCACCA CGCCAGGCTA
75541   ATTTTTGTAT TTTTTTTAGT AGAGATGGGG TTTCACCATG CTGGCCAGGC TGGTCTCGAA
75601   CTCCTGACCT CAGGCGATCT GCCCGCCTCA GCCTCCCAAA GTGCTAGGAT TACAGGCGTG
75661   AGCCACCAAG CCTGGCCTAA GTGACATGTT CTTATATTGT TCCTTTCTTT CTTTTTTTTT
75721   CGACTGAGTC TCACCCTGTT GCACAGGCTG GAGTGCAGTG GCGTCATTTC GGCTCATTGC
75781   AACCTCTGCT TCCCGGGTTC AAGCGATTCC CTTGCCTCAG CCTCCTGAGT GCCACCACCC
75841   CCAGCTAATT TTTGTACTTT TAGTAGAGAT GGTGTTTCAC CATGTCGGCT AGGCTGATCT
75901   CAAACTCCTG GCCTCAGGTG ATCCGCCCCC GAGTCTCCCA AAGTGCTAGG ATTACAGGCG
75961   TGGGCCACGG GGCCCAGCCT TATATTATTT CTTTTACTAC AATATATTAG TATGATGCAG
76021   GTGCTTCAAT TGTTTATACA CTTTCCATAA TTTTGTATAA TTCTTATACC CTGTCACTCT
76081   GAGGAATAGC CGGTCTAAGT GTTTTTCCAC CACTGCTAAT TCATCCATCA CTAATCTCAT
76141   TAGACTGTTA ATTCCCAGAG GACATAAGCA CACAAGCAGA CAATGTTTAC AAATGTTGGA
76201   CAAATGTTAT TTAATAAAAC AATGGGGTCA CCCTTAGTCT AAAAGATGTT TCACTTTTCA
76261   TTTGTCATTG AACTCTTATT TGTAGGTTCC CTTTTGACTT TCCCACAATC TAAGGCTGTT
76321   CTCTTTAACA CATATTTTCA TGAAAACATA TATTTGAGCA GAAATTGTTG GGGAGTTGTA
76381   ATATTACCTT TGTCCCTAAA TATGAATCTA TAATTATATC AAATATATGG GCAGACAATT
76441   TACTTTGCCT TTAATCTCAA GAAAAAAATA GCAATTACTT GGGGTCGGAG AGTAAAATAA
76501   GAAGTAGTGA ACCTTAAAGT AGCAAACTTT AGAACAGAAT AGTTTCAGAG GGGATGAGAA
76561   GAGGTGATTT TTCAGCTCAT CAACAACAGA TCTTATAATA AATTACATGT TCTGGTACTT
76621   TTCTTGTCTT TCTGTGTTAA ATTTTGCTAT TTAAAAAAAT AAATTTCAAA TACATTGTTC
76681   ATCTTAAAAG TCAAGAGTGT GTTTTATTAA AGTCAGTTGC TTTATTTGCA ACTCAAAAGA
76741   TATATTTGAG TTCCCAACTG GAGATTGTCC TATATGGTAA CTTGCGTAAG GTATGGTTAC
76801   TGAAAGTAAC CTACAATTTT CATGGGCTGA AATTCATTTC TATATTGCAG CGTACAAAAA
76861   TAAATAAATA AAAAATGCTT GTTTTCTTTG AAAACATATT ATCTCAGTGC CTCTAACTGC
76921   CAAATCTATT GGCTTTTTTG CAGGCTTAAG GGCTCTCCCT TGTTCCTTTA TGATCTCTAT
76981   CTTGAGGGCC AGACCTCCTG CCTTACACAA CTCAGAGGGG GACCTCAGAG CTCTTTAAAA
77041   AGAGCCCAAT TTCTCGCCTG TAGAGAAGTG AAAAGGATGC CCCACCCCCA TCTATGAAAA
77101   GAGGGATTTG ATAGTTTCAA TGTCTTCAAA TCAAAGATTT AAGTCTGTAG CCCCCCACCA
77161   CCCCGGACCC TAGCAAGGCT CATGAACCCC CTCCCATCCC GCCCTAATTG CTTTGGACTG
77221   GCCGTGGAAT CCTTGTCCCA GTCCACAGTT CCTGTGCGAC TGCACGAAGA ATTCACAGAG
77281   GACCTGTGTT ACTTCCCTTG TGAAGAAACA GAATTATCAT GAAAATTTAG GTGGAAACCA
77341   TTTCGCTTTT TTCTTCAAAA ATAAGGGAAG CATGTGCCCA ACCACCCCTG GGAAAAAGAA
77401   CCTTCAGGGG CAAAGGAGCG AACAGGTAAT TTATAAGAAA AACAGAAAGT GGTCTCTGAC
77461   TGCCCCAGAC TTCCTTCGGA GTTGGGGGAA TTGGGGACGC CTGGACGCGT TGTTTTTGTG
77521   TTTGTGGAAA AAATAAATGA AGAGCATGAA GCCCGAGGCT TCTGAGATCC TTTCCTGACC
77581   AAACCCAAGT GATTTGGTGC GGGGAATTTT AATATTTTTC CCCTTTTGTG AGGTGGAACA
77641   AACACAACTT GGGAGCAGCG CAGCGGCTCA GAGCCTGCCA GCCAGGCGGG CGACCAGAGC
77701   ACCAATCAGA GCGCGCCTGC GCTCTATATA TACAGCGGCC CTGCCCAGGC GCTGCTTCAT
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77761   CGGCGCTTTG  CCACTTGTAC  CCGAGTTTTT  GATTCTCAAC  ATGTCCGAGA  CTGCTCCTGC
77821   CGCTCCCGCT  GCCGCGCCTC  CTGCGGAGAA  GGCCCCTGTA  AAGAAGAAGG  CGGCCAAAAA
77881   GGCTGGGGGT  ACGCCTCGTA  AGGCGTCTGG  TCCCCCGGTG  TCAGAGCTCA  TCACCAAGGC
77941   TGTGGCCGCC  TCTAAAGAGC  GTAGCGGAGT  TTCTCTGGCT  GCTCTGAAAA  AAGCGTTGGC
78001   TGCCGCCGGC  TATGATGTGG  AGAAAAACAA  CAGCCGTATC  AAACTTGGTC  TCAAGAGCCT
78061   GGTGAGCAAG  GGCACTCTGG  TGCAAACGAA  AGGCACCGGT  GCTTCTGGCT  CCTTTAAACT
78121   CAACAAGAAG  GCAGCCTCCG  GGGAAGCCAA  GCCCAAGGTT  AAAAAGGCGG  CGGAACCAA
78181   ACCTAAGAAG  CCAGTTGGGG  CAGCCAAGAA  GCCCAAGAAG  GCGGCTGGCG  GCGCAACTCC
78241   GAAGAAGAGC  GCTAAGAAAA  CACCGAAGAA  AGCGAAGAAG  CCGGCCGCGG  CCACTGTAAC
78301   CAAGAAAGTG  GCTAAGAGCC  CAAAGAAGGC  CAAGGTTGCG  AAGCCCAAGA  AAGCTGCCAA
78361   AAGTGCTGCT  AAGGCTGTGA  AGCCCAAGGC  CGCTAAGCCC  AAGGTTGTCA  AGCCTAAGAA
78421   GGCGGCGCCC  AAGAAGAAAT  AGGCGAACGC  CTACTTCTAA  AACCCAAAAG  CTCTTTTCA
78481   GAGCCACCAC  TGATCTCAAT  AAAAGAGCTG  GATAATTTCT  TTACTATCTG  CCTTTTCTTG
78541   TTCTGCCCTG  TTACTTAAGG  TTAGTCGTAT  GGGAGTTACT  GAGGTATCAG  ACGAATTGGG
78601   TGACGGGGTT  GGAGAGTGGC  CGTGGTGAGG  TTACAGCATT  TAAACCTTTA  TTGCGGCTTC
78661   TAGGTCCCTG  ACCGGAGGCT  TTTCTCGCTG  GCGGATGGTT  TTGGGATGGC  AGTCCCGCCC
78721   CAGGCCTGTG  AACGGCAGAA  AAGACCGCAA  AACAAGAGCC  AGTTTCTTAG  TCTAAAGGGA
78781   TGTCCGGATT  GGACTAAAAA  ATTTTCAAAA  GTCCCGCCCT  GCTCCCGGGT  TGGTCCGTTC
78841   TTCTAGTACA  TGACTTTCAT  TCTGTATTTA  ATTGGATGGT  GGAAGACGTT  GCTTATTCTG
78901   TGTTTTTTGC  TTTACTGTGA  CTTAAAAGTT  TTGCCTCTTT  TCTCTTTATA  TTAATGTCTG
78961   GGATTTCGGA  CGCTTTCCAT  GTTGTTGGTA  GTCAAGTTGA  TGTCTCCTGG  AGGTAGTGGC
79021   AACATCCAGC  CCTGGGAGGA  GAGTGCGTGC  AGGTACCTTT  GTCCTACATT  CCTCTGCTGT
79081   TAATTTCTCA  TTCCTGTGGC  AACGAAGGAA  TGCATTTAAA  AAACAGCCAC  AACAGCGGCA
79141   ATAGCCCTTC  CTCCACCCAA  GGCAATCGTG  GACCTAGGGA  GTTTTTTGTG  CCACATAACA
79201   TGTAGCCTTC  CGCTAAACTG  ACAGGTTTGA  GCGTATCGAT  TTTGAGCGTA  TCGAAAGCAC
79261   AACTTTTAGC  CAGCCATTTT  GTCCTCGCAT  GACTACGGTT  GCTTATCCTG  TTTAGACAGA
79321   CAGCAACATT  TAAAAATCGA  AGTTCCTTTA  AACGTATTTT  GTTGGCAGT   CCAAATGTTT
79381   CTATGCAGAA  AACAGTATTT  GTACTATTAA  CTATGAAGAG  TGTATGGATA  AATGGGAGAC
79441   ATTTCTAATA  AAGGCCTTCG  TTAATGGTTC  CCTCTGTTTG  ACATCCATGG  TGCTTCTGAA
79501   TACAGAAAGC  CTAGCGTCTT  ATATTCGCTT  CTTTTAAAAT  CTGGTGGGCA  CATTTTGGTG
79561   AGACCTAAAT  TATGGGGACT  GGGGCTTCTG  GAGATAAGCT  GCTCAATTAT  TCTACCATCT
79621   CCACAATGAT  TAATATAGTG  AGTTGATTTG  TTAGTGATAG  TGACCACGGA  TTCATCCCAA
79681   GAAAGAGAAA  GGGGAGGGAG  GCAAGCAGAG  AGACAGGAAG  ACAGAGGCAG  GGAAGAAGGA
79741   GAAAACATTC  TCCCATGGTT  TAAGTAATTT  TGTGTTGTTA  ATTTTACATT  ACAACACGGT
79801   TTAACATGGT  GAACCCTCTA  TTTTGGTGTA  AGGTTTAACA  TATGGACATA  TTTTTCCCAA
79861   GACCATTTAT  GAACTTTCAT  TTCTGCTTCC  CCCTTCTTCC  TCCCGTGCCA  CCCTCCACGC
79921   TCCTATCAAT  TTTGGCTGTT  TTGTCATAGG  CTAATACGCT  ATAATTTCAT  GGACAGTTGG
79981   ACTGTCTTAG  GTTTCTCAGG  TTTCTATTTT  GTTCCTTTAG  TCATTCCCAC  AATTCTTAAG
80041   GTAGAATTGT  ATTGTTTTAA  ACATTGTGTT  GTGTGCTATC  CTCAATGCTG  AGATGATTAT
80101   GTGACAAATG  GCAAGTGTTC  AACTAATACC  TAAATCTGTA  GTATCTTATC  AAGCCTAATG
80161   CTACTTCACA  ATGCCTACTC  CATTCACCTC  ACTTTATCTC  ATTACTGGCA  TTCTGTCATC
80221   TCACATCATC  ACAAGTAAAA  CGGTAAGCTA  TTTTGAGAGA  GATCACAGTC  ATATAATTTA
80281   TATTTATATT  TATTTATTTA  TTTATGAGAC  GGAGTTTCCC  TCTGTCACCC  AGGCTGGAGT
80341   GCTGTGGCAC  GTTCTCGGCT  CACTGCAACC  TCCGCCTCAC  GGGTTCAAGC  GATTCTCCTG
80401   CCTCCGCCTC  CCGAGTAGCT  GAGATTACAG  GGGCCTGCCA  CCATGCCCGG  CTAATTTTTG
80461   TATTTTTAGT  AGAGACGGGG  TTTCACTAAG  TTGGCCAGGC  TGGTCTCGAA  CTCCTGACCT
80521   CAGGTTATCC  GCCCACCTCA  TCCTGCCAAA  GTGCTTAGAT  TACAGGCGTG  AACCACCGTT
80581   CACAGACTCA  AATCATTTTT  ATTACAGTAT  ATTGTTATAA  TTGTTGTTTT  ATTATCAGTT
80641   ATTGCTAATC  TCTTACAGTG  CCTGATTTAT  AAATTAAATT  CATCATTGCC  ATGTGTATAT
80701   AGAAAAAAC   AGTGTATATA  CGGTTCAGTA  CTATCTGTGG  TTTCAGGCAT  CCACTGGGGG
80761   TGCAGTTTAT  TAAACATGCA  TTTACATTAG  TCTCCCCTTT  GGGAGACTAA  TTAACTGAGA
80821   TGTTGTAACG  TGACTTTAAT  AGCAGATAGA  GCTAATTTTC  TCTCATTACT  CTTCTTTTTC
80881   AGAATTTTCC  TGGTTATTCC  ATTTTTTATT  TTTCCATATG  TATATTAAGA  TCTCTTCCAC
80941   CTCCTCCTGT  TTCTCCATCT  CAACATCAAA  CAATTAAAAA  AAAAAAAAAG  GCTGGGCGCG
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81001  GTGGCTCACG CCTATAATCC CAGCTCTTTG GGAGGCCTAG GCGGGTGGAT CACGAGGTCA
81061  GGAGTTCAAG ACCAGCCTCG CCAAGATGGT GAAATCCCGT CTCTACTAAA AGTATAAAAA
81121  TTAGCCAACC ATGGTGGCAG GCGCCTGTAA TCCCGGCTAC TCGGGAGGCT GAGGCAGAGA
81181  ATTGCTTGAA CCTGGGAGGC GGAGGTTGCA GTGAGGCGAG ACCTTGCACT CCAGCCTGGG
81241  TGACACAGCG AGACTCCGTC ATAAAAAAAA AAAGCCGGAA GCAGTGGCTC ACGCCTGTAA
81301  TTCCAGCACT TTGGGAGGCT GAGTCAGGCA GATTACCTGA GGTCAGGAGT TCAGGACCAG
81361  CCTGGCCATG AAAATACAGC CTGGCCATGA AAACACACAA TAAATTAGCT GGGCGTGGTG
81421  TCACACACCT GTAATCCTAG CTACTCGGGA GGCTGAGACA GGAGAATCAC TTGAACCCAG
81481  GAGGCAGAGG TTGCAGTGAG TTAAGATGAC GCCACTGCAC TCCATCTGGG CGACAGAGCC
81541  AGACTCTCTC TCAAAAAACT AAATAAATAA AAATAAAGTT ATGGTACATT GAACTTCTGT
81601  GTTCCTTTCT CCCTTAGATA CTTTCATGGC TACCCATTTA ATTGATGTTC TTATCATCTC
81661  CAAGAGTTAG TCAGGAGAGG AATCAACCCA AGCAAAAATA GCTGATTTTC TAATTTTCCT
81721  TCAATGCCCT TTGGGGTCTT AATCCATTTG ATTTATGTAC TTTCAATTAA TCCTAACCTC
81781  GAATGTCTTC TGCAAACATG TTTCCACAGA TGAAACTCGT CAAATGAAAC ACATTCCTTT
81841  AATTTATAGA GTTAAAAATT AGAAAATTT  TCAATTCTAT TTGGCCTTTA GATTCAGTCT
81901  TGCATATGTT TTCTCAATTT TGTTCATGCT CTTTAGTTTT GTTTTATTCC ATCACAATTG
81961  TTCACATAGC TTACTGGCTT AGGTCTAATG AACCATTCAT TTGGAAATTA AAATTGGCCA
82021  TTTTAAGATG AAAAAGATTC TTGCCTCAAT TTTACTTAGT TTTTGAAACT GTCAATGAGG
82081  ACACATGTTT TTCTGTACTC TTAGATTCAC TAAGTAGTGT CTTGCAAATT TAACTGACAA
82141  AGGACAGATT AACATGCGAA AAAAAGAGCA TGCAATTTTA TTAGTATATT ACATGCACAG
82201  AGTTCCCAAA GAAAAAAAAA TTGAAACCTT AAAAACGCGG TTAGACTCAC AGACTTATAC
82261  ACCATTCCAA CAAAGGAAAG GGAGTTTGCA CTTCATGGGA TGACGAATTT GGGAATGTGA
82321  CAAGGAAATA AATACATGGG CAATAAAAAC CATGGAAGAT AAAATGAAAG ATAGAAATAA
82381  TTGTAGTAAG GTTTGTTTTT GCAGAGTCAT CTCAGTGCCA ACCTTCCATA TCTAGTGATA
82441  AGAATTGCTC TCTTTTTCCT GGTATAGCAG TTGGGGACAC TTTTACAAGG GAAATTTCTG
82501  TCACCTTCAC AAAGGGAAAT TTGGGTAAAG AGAAGACAGA GACCTCTTCC TACACCTGTT
82561  GATTTTCAAT TGCCTTCAGC TGAAAATAAC TTTTATGCCA AAGTAGAATA ATTTGGGGGT
82621  GACATCCTGA TATTCTTCAA AACTTATATT TAATTTCACA TTAGTAATTA TATCATTTTT
82681  GATTTTTAAA TTAGTTTTAT AAAATAATTT TGAAAAACGG TAATAATATT CAAATAATTC
82741  CAGAAACACT GCTGATAAGC CAAAAACATC AATGAATATT GCATAAACAA CTGATAATTC
82801  AACCATGAAA ATTTATGACA TTGTTCTTGT GTGATAAAAC TATGAGTAAC ATAAAAACTA
82861  GAGGCTACTT GTAATGCATT ATTCCAAACT TTCTGTTTTT TATTTATTTA TTTATTTATT
82921  TTGAGACATA GTCTCTCTCT GTCACCCAGG TTGGAGTGCA ATGGCGTGAT CTTGGTTCAC
82981  TGCAGCCTCC ACTTCCCCGG TTCAAGCAAT TCTCCTGCCT CAGCCTCCTG AGTAACTGGG
83041  ATTACAGGCA CCTGACACCA AACCCGGCTA ATTTTTTTGT ATTTTTAGTA GAGACGGGGT
83101  TTCGCCATGT TTGCCAGGCT AGTCTCGAAC TCCTGACCTC AGTGATCCAC CTACCTCGGC
83161  CTCCCAAAGT GCTAGGATTA CAGGCGTGAG CCACCATGCC CGGCGCATTA TTCCAAACTT
83221  TCATACACAG TGCTATCATG GCTACAAATT GAAGTATCAT ATTATACACT CCTAGGCAAA
83281  GCTCTGGATA TTTTGGCTAT ATAAGCCTGA GGGAAATGTA GTAAGGACAT TGTGGTTGAA
83341  ATTCATACCA GAGATGAACA GGCCCAGTGC AAGACAGAAT TACATCACTA AAGGATATCA
83401  GAAGAGAATA GGGATTTAGG GTACAGTGGC AACAACAGTT TTGGGAACTA GCATTTTTTG
83461  AGCACTTATT TACAATATGC CAAGCACTGT TGCTGATTAC TCTATATTTA TTTTCAAACA
83521  CATTCTTGTC ACAGCACTTT GAAGTAAGTG CCATTGTCAT TCCCACTTCA GGGTGAAGGA
83581  CTAAAGCTTG GTGTCATTAA GGATGTAGCT AGTTAGCTGT GTGTGTGTGT GTGTGTGTGT
83641  GTGCATTTTT TTTTAAATTT AAAGTCAATA AATTTTTATT TGAAGAATTT CACATCAAGG
83701  TAAACTTTGT TCCTCTAAAG AGCTGGAGTC AAAATGTATC TTCAAAAGAT TCATCTTCAA
83761  GTTAGCCCTT CTTAATAGAA CTGATGCTTA ATCCACAGTT GTCAGCCCAC AGTTCTTTTA
83821  TTTTGACTTT TTTTTTTTTT TTTTTTGAG ACGGAGTCTC TCACTGTCAC CCAGGCTGCT
83881  GGGCAGTGGC GTGATCTCGG CTCGCTGCAA CCTCTGCCTC CCGGGTTCAA GTGATTCTCC
83941  TGCCTCAGCC TCCTTAGTAG CTGGGACCAC AGGCGCATGC CATCGTGCTC GGCTAATTTT
84001  TGTATTTTTA TTAGAGACAG GGTTTCACTA TGTTGGCCAG GCTGATCTCA AACTCCTGAC
84061  CTCATGATCC GCCTGCCTTG GCCTCTCAAA GTGCTGGGAT TACAGGTGTG AGCCACTGCA
84121  CCCGGCCTTA TTTTGCCTTC TTTAATCTCC ATTTGAACAT ACACATACTG ATGAAAACTA
84181  CAACATTCTT CACCAAAAAT CTTTGGGATT TAATTTCTTC AACCACTTTA CTTTGGGGTC
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84241    ATTTTAAGAT TAGGTGTATC TGCCTGGTTC TCAATTTGAC ACCCTTTCTC TCTAAACATG
84301    AATGAGTTCC AATCATATTT ATTCCTAAGC TATCACACTC AAATATACTA CAGATCTGTG
84361    GAATATGCCA AAAGTTAAGG TGAAAAATTA AATTATTAGG TATTTCATAG TTTTGCTAGT
84421    TTTTGATCTG TGAGTGAATA TAACTATCCT CTATGTCCTG GCACTGTTCC TCAGAAACAT
84481    AGGGTCCACA TATGTAATTT TAAATTTTTT AATAGGCACA TTTTAAAAAG TGAAAAAAGA
84541    AATCTATTTT AATGATTTGA ATCCAGTGTA ACCAAAAATT GTTTCAACAA GGTATCTAAT
84601    ATTAAAATAT TGAGTTTTTA CTTTGTTATT TTACTAGTTC TTTGAAATCT GGTGTGTATT
84661    TTACACTTAA AGCACATCAC AGTTTGGAGT AGCCACATTT CCAATGCTTA ATACTCACAT
84721    ATGGTTAGTG GCAACTATCT TGGACAGGAC AGCTTTTATA CTCTGGGAAG ACACAAGCAA
84781    ATACTTGCTC TGCAGCAGAA TCCAGATGTT TTCCAAGAAA ACACTTTTTC TGACCTGTTC
84841    CTGAAACCCA GGTAGTGTCT CTAATACTTT ATATTTTATT GGTTGTCCT ATTGTAACCA
84901    CCCAACGGGC TCTCCTTGTC CACTTCCTAG ACAGAGCTGA TTTATCAAGA CAGGGGAATT
84961    GCAATAAGGA GCCAGCGCTA CAGGAGACTA GAGTTTTATT ATTACTCAAA TCAGTCTCCT
85021    TGAGAATTTG GGGACCAAAG TTTTTAAGGA TAATTTGATT GTAGGGGACC AGTGAGTCGG
85081    GAGTGCTGCT TGGTTGGGTC AGAGATGAAA TTATAGGGAG CCTAAGCTGT CCTCTTGTGC
85141    TAAATCAGTT CCTGGGAGTG GTGGGGTGGG GGACTCAAGA CCAGATAATC CAGTTTATCT
85201    ATATGGGTGG TGCCAGCTAA TCCATTGTGT TCAGGGTCTG CAAAATAGCT CAAGCATTGA
85261    TCTTAGGTTT TAAAATAGTG ATTTTATCCC CAGGAGCAAT TTGAGGTTTA GAATCTTGTA
85321    GCTTCCAGCT GCATGACTCC TAAACCATAA TTTATAATCT TGTGGCTAAT TTGTTAGTCC
85381    TGCAAAGCA GTCTGGTCCC CAGGCAGGAA AGGGGTTTGT TTCTGAAAGG GCTGTTATTG
85441    TTTTTGTTTA AAAGCAAAAG TATAAACTAA GCTCCTCCCA AAGTTAGTTA ATCCCAAACT
85501    CAGGAATGAA AAGGACAGCT TGGAGTTTAG ACGTTAGATG GAGTCGGTTA GGTAAGATCT
85561    CTTTCACTGT AATAATTTTC TCAGTTATGA TTTTTGCAAA GGCAGTTTCA CTGTCCACTT
85621    CACCTCACAT CAGGCCTCTG ACTAGAGGAT TCCAACAATA CTTAGGCCAG GACACCACCA
85681    TGTCTCCTTA TCCACCCTGA GGGAGTCCAA TTTCTGAAAC AAAGGAAACT ATATATGATA
85741    GTATGAAACT ATATATGAGA AGGAAATTAT ATATGATAAT CAATTTTAGG GTTATCTTAT
85801    TGATTAGAAG ATATTAAAGT GTGACACTGC CTGGCAATGA TATCTGCTGG TAGTAAGAAT
85861    TTGGCGAATT TAGTGAAATT CCTGAGGCTG AACCTCCACT TCTGTAAAAT GGAGACAGTG
85921    AGATAATTTG CCTTACAATG CTGAAGTAAG AATTTTACAC AATAATTCAG ACCAACCACT
85981    TCATGTGGTA CTTGGCCCGT GGAAGACTAT CAATGACAGT TAGTTTATAG TTTATACTAT
86041    TAATGAATCC TTTGTTTCAT TGTTATTTCC TTCTACACGT TGGCCTCTCT AAAAGAAGGT
86101    AATATTCAAT ACAAATAAAG TTAAAACAGC TTGCAGAGTT GTCCCAGGGA ACTCACTTAA
86161    CCACTGAAGT GTTCAAATTG CTTAAGGTTG ACTTTATATT CTCCTGACTA ACCTTTCTCC
86221    TTCTGGTATT TCTTCTGAGA ACAGCACCAC CATCCAAAGC ATCATGCAAA CAGTGGTCAT
86281    CCCAGACCAG TAATTCTCAA CTCACAGGGT GCTCCTGCAG AGATGTATTT GAATAGAGTG
86341    GTAGGATGCT GAAGAAGGCC ACGTAAAATT TGGCCAGTGA TCTGGGGCAG ATTTATCCTG
86401    AAGCTAATGA AACACAAGTG TAAGGGCCTG TACTTCCAAG GTGCAGAGAG GGGCCCTACA
86461    AATGTGTTAG TTTGTCTCTC TCTCTCTCTC TGATTTTAAA ATTTGCAGTA TTAAGGTACT
86521    TTAATCACGG ATGGTTCAGG CTGCTATTTT CACTCAATCC TCCTTTTTAT TAAAATCACC
86581    ATTGTCTGAT TATGTTAGAA TCCTGATGAA AATATTTGGA ATTTGAGTAA GAGAAAGTTT
86641    AGTTGAAGAT GTATCTAGTA TGGGGATAAT AAGTTACGTG ATTTGCATAT GTGATCATGT
86701    GTACTTCATT CGTTGCCAGC CAATCTGACG TAAGAATGGC TTCAAGGAGG CCGGGCGCGG
86761    TGGCTCACGC CTGTAATCCT AGCACTTTGG GAGGCCGAGA CGGGCGGATC ACGAGGTCAG
86821    GAGATCGAGA CCATCTTGGC TAACACGGTG AAACCCCGTT CTACTAAAA ATACAAAAAA
86881    TTAGCCGGGC GTGTTGGCGG CGCCTGTAG TCCCAGCTAC TTGGGAGGCT GAGGCAGGAG
86941    AATGGCATGA ACCTGGGAGG CGGAGCTTGC AGTGAGCCGA GATTGCGCCA CTGCACTCCA
87001    ACCTGGGAGA CACAGCGAGA CTCCGTCTCA AAAAAAAAAA AAAAGAATG GCTTCAAGGA
87061    ATGTTCCTAC TGCTCACTGG AATAACTCAC CTAAATTCCT GGCAAGATGC AGGTCTAGAT
87121    AAAATGTTAT GACATCTAAG TATTCAAAAC ACATTCCCAG CACTGAGAGT GAGTGTCTAG
87181    TGGAGAGTAG AAACGTATAG AGCCAGAAGC TAGTCTGGAA AGAATTCTTA CAAAGTTTAC
87241    AACTTACATG TGAAAGGAGC TTAACAGAGG ATTTTCCAAA TTTGAAAACA ATCCTAAAAA
87301    CTTACTTGAC ATTACCAATA ATGTGTTTTG AAACTGAAAT ACTTCTAAGT TATGAAGAAA
87361    ACATATTATC ATCAGCCACC CTGGAGGAAA GATTGAATTC TATTTCCATT ACCTATAGAC
87421    AACATTACAA AATAATTTCG ATCTGAAGAT GGAATCAGAG TATTCAGTCA AAACTACAGG
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87481   AAAATATACT TGGTAGTGTC ATATTCAGAA GTTAATAAAA TATGCTATTT TCTGAATTTT
87541   GTGATGGCTG TTGTTTTGTC AGCTTTTATA AAATTGGAAT TTGATTTTAT TTTCCCATTA
87601   TAAATTTATA TTTACAGTCT GCAGTACTTT TGCATTTTTA ATTTTACATT ATAGTTTTTA
87661   ATAGTTAACA AGTTGTAAAA GGTTTGATCC CCAGAAAACC TTGATCTACC CCATCAGTTA
87721   AGTATACTAA TATATTTAGA AAATGGATGA AATCAGCATT TGAATATTTT TAAATATTTA .
87781   TTAAAGAGG ACATGGGTAA AAGAGCTTTG CAGTTGCCAC CCTTCATTCT CAAATTCCCT
87841   GGATAAGGAT GACCGCATAA TCTTTGGATG GTCATACGCA AGTCTTGTGT ACTTGTTACA
87901   TAAATCTATT TAGTGGACTT TTGGCAGTGT GTACTGAGGC CAGTTTCTTC CACCTGAGCT
87961   CTGACTCCAC CTCCAGCAGC CCAAAACCAA TACTGAATTT TGGGGTCAGC TATTGTTTTT
88021   GTGGACTTAG GTAACTACAC ACACATTGTC TTTATGATAG CTTTAATAAT ACTGCCATCA
88081   GAACTAAAAT TGTCACGTGG ATTAAAAGGA GTGACGGTGG TGTCCCCAGG AGCCTTTCAA
88141   TATGTAAGTA TTTACACATA TACATGCTAA AAAGACCCCT AGGAATTTTT TAACAAGGGC
88201   AAAACAGTAA CTCAGCTTGT TTTCTCGCAG TAAAACCGGT TGAAAAGGCC TGATAGACTT
88261   GTCTGCAGTT ACAAAACTTG TGTGTAGTTA TCACCTTTAT ATCTCCTGGA AACTAACATA
88321   GACAACCGAA TGGGTTACAA CTGTTTTTAA GTGAAATTGT GAGTGGCTCT GAAAAGAGCC
88381   TTTTCAATGA GGAAGAAACG GGCAGACTTA TGCCCTTTCC CCACGGATGC GACGTGCCAG
88441   CTGGATATCT TTGGGCATGA TGGTGACGCG TTTAGCGTGA ATAGCGCACA GATTGGTGTC
88501   TTCGAAGAGT CCCACCAGGT AGGCCTCACA AGCCTCCTGC AGCGCCATCA CCGCAGAGCT
88561   CTGGAAACGC AGGTCGGTTT TGAAGTCCTG GGCGATTTCT CGCACCAGGC GCTGGAACGG
88621   CAGCTTCCGG ATCAGCAGCT CGGTGGACTT CTGGTAGCGA CGGATTTCGC GCAAGGCCAC
88681   GGTGCCCGGG CGGTAGCGAT GAGGTTTCTT CACGCCACCG GTGGCCGGAG CGCTCTTACG
88741   GGCTGCTTTA GTAGCAAGCT GCTTGCGCGG AGCTTTGCCG CCGGTAGACT TGCGAGCTGT
88801   TTGCTTCGTA CGAGCCATTT GCAATGAGAG CACACACAAA AGTGTAGTGA ACTGAGAGCA
88861   AGTGGCCTTT AAATATAGTG AGAAACATTC TGATTGGTCC TGTAATATTT CAAAAGTCCC
88921   GCGCGATAAA ATCATTGGCT GAAGAGTGAC CAGACTGATT GGTTCATTAC TAGACAATCT
88981   TATTGGATGA GTTGCCCCAC CGCCCATCCT GTCCTTTTCG TTTCAGTTAT CTGCAGCGAC
89041   AAATTGTCTA AAATTCTAGT TCATCCAGTC CCAAAGAACA GAGTGTATAA CAAGGTATCT
89101   AAGGATTTTT AAAATGTAAA TTCCGATTCA GTAAGTTTGA GTGGGACTTG AAATTCTGCA
89161   TTCCTGACAG TCTCGCAAGT TATCAATGCT GGTGAACACT CACTAAACCA CCAGAAACGT
89221   TCAGACTCAT GTCGGGAAAT AACGCTTATA TTCAGAGAAT GAGATTCCAT GCTATTTTGT
89281   TACTGGCGAA CAGCAAGTTT CCTTGCCCTT TGTTTTCTAA GTCCAAGTCA CATTCCCACC
89341   CTGCCTGTTC TCAAAATGTC TTATTTTGGT TGGCCTTAAG TTTCACTTTG TATACTCTAA
89401   AATGTACTTT CTAAAGGAAG GTGTTATTTT CTCGAAACTT AACTTTTTAA CACCATTAGG
89461   CTAGGGGGGC GGTGGCTCAC GCCTGTAATC CCAGCATTTT GGGAGGGCGA GATGGGACGA
89521   TCACTAGAGG CCAGGAGTTC AAGACAACCC TGGCTAAAAT GGTGAAACCC CGTCTCGCAT
89581   AAAAATACAA AAACTAGCTG GGCGCGGTAG CAGACGCCTG TAATCCCAAG TACACAGGAG
89641   GCTGAGGCAT GAGAACCGCG TGAAGCGGCG GGGTGGAGGT TGCAGTAAGC CGATATCGCG
89701   CCGCTGCACT CCAGCCTGGG TGACAGAACT AGACTGTCTC AAAACAAACC AATCCAAACG
89761   AAAAGCAAAA AATACCCTAA CAGAAGCAAG TTATCATCCT TTCTTGTGTA ACTATGGACG
89821   GCTCTGAAAA ATGCCGTTTC AAGTGTAAGC TACGTTTTCT GATTTGAGTG TTTACTTGAC
89881   CTTGGCCTTA TCGTGGCTCT GTTATTTTGG CAACAGGACG GCCTGAATAT TGGACAGGAC
89941   GCCTCCCTGA GCAATAGTGA CGTTGCCCAG CTGCTTGTTG ACCTCCTCGT CGTTTCGGAT
90001   GGCCAGCTGC AGGTGGCGGG GGATGATGCT GCGGGTCTTG TCACGTATGG CGCTGCCCAC
90061   CAGTTCTAAG ATCTCGGCGG CCAGGTATTG TAAGTACACT GGCGCACCGG CTCCGACCGG
90121   CTCAAAATAA TTGCCCTTTC GAAAAGATG ACGGACTCTG CCCTATTGGG AACTGCAAGC
90181   CCGGTAGCGA CGAACAAGTT TTTGCTTTAG CTCCATTTTC CACGTCCGCA AATAGCGACC
90241   TATGAAAGCA GCGGAAAACT GTGAAAGACA AGCAAGCTGG AATGGCGCCT GAACAAATCC
90301   TTTTATACAA ACTGCAAGGC TGCAATAGGA AGCTATCCTA TTGGTCAATT ATGTTTGGTG
90361   CTTTATCCAA TAGAAAAGA TAACATAAAT TCCATATTTG CATAAACCCC ACCCCTCAGT
90421   GAAACCGTGT TTCTTTTGTC CAATCAGAAG TGAGGAATCT TAAACCGTCA TTTGAATCTC
90481   AGGACTATAA ATACATGGGC TCTGAACTGT TCTCTGTACT ACTCTGTAGT GGAGAGTGTT
90541   AGTAGCTTTT CTATTCTGTT TAGGAATAGC AATGCCTGAA CCCTCTAAGT CTGCTCCAGC
90601   CCCTAAAAAG GGTTCTAAGA AGGCTATCAC TAAGGCGCAG AAGAAGGATG GTAAGAAGCG
90661   TAAGCGCAGC CGCAAGGAGA GCTATTCTAT CTATGTGTAC AAGGTTCTGA AGCAGGTCCA
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90721    CCCCGACACC  GGCATCTCAT  CCAAGGCCAT  GGGGATCATG  AATTCCTTCG  TCAACGACAT
90781    CTTCGAGCGC  ATCGCGGGCG  AGGCTTCTCG  CCTGGCTCAC  TACAATAAGC  GCTCGACCAT
90841    CACCTCCAGG  GAGATTCAGA  CGGCTGTGCG  CCTGCTGCTG  CCTGGGGAGC  TGGCTAAGCA
90901    TGCTGTGTCC  GAGGGCACTA  AGGCAGTTAC  CAAGTACACT  AGCTCTAAAT  AAGTGCTTAT
90961    GTAAGCACTT  CCAAACCCAA  AGGCTCTTTT  CAGAGCCACC  TACTTTGTCA  CAAGGAGAGC
91021    TATAACCACA  ATTTCTTAAG  GTGGTGCTGC  TGCTATTCTG  TTTCAGTTCT  AGAGGATCAA
91081    CTGGAATGTT  AGCGAAGACA  AGTTTTAGAG  CCAAGGTTAA  CTTGGACGGG  GCCGTGCGCG
91141    GTGCCTCTTG  CCTTTAATCC  CGGCAATTTG  GGAGGCCGAG  GCGGGCGGAT  CACGAGGTCA
91201    GGAGATGGAG  ACCATCCTGC  TTAACACGAT  GAAACCCCGT  CTCTACTAAA  AATACAAAAT
91261    AATTAGCTGG  GCGTGATGGT  GGGCGCCTGT  AGTCCCAGCT  ACTCGGGAGG  CTGAGGCAGG
91321    AGAATGGCGT  GAACGCGGGA  GGCGGAGCTT  GCAGTGAGCC  GAGATCGCGC  CATGGCACTC
91381    CAGCCTGGGT  GACAGAGCGA  GACTCCGTCT  CAAAAAAAAA  AAAAAAAAAA  AATTAAAAAA
91441    ATATGAAGTT  TTGAAGCAGA  AATTATTTTG  TCGTATGTTC  TTTCATAAAT  TTTTTGCCTG
91501    CCTGCCTTCT  TCCTTTGTTA  CAGAACTCCA  ACACTTACCC  AAAGGTAGCT  GTTGGGTCAG
91561    GGTTTCTGTA  CTATAGTCCC  TTCTGTGGTG  GCCAGAAATA  TGTTACAGGA  AAGAGGTCCC
91621    CATCCAGACC  CCAAGAGAGG  GTTCTTGGAT  CCCGCGCAAG  AAAGAGTTCA  GGGTGAGTCC
91681    GCAGTGCAAA  GTAAATGCAA  GTTACTAAG   AAAGTAAAGT  GGTGAAACGA  CAACTACTCC
91741    ATAGACGGAG  CAGGACATTC  CCGAAAGTAA  GAGGAGGAAG  GCATCCACCC  TAGGTACAAT
91801    ACTTGTATAT  ATGGGGAGAT  GTGCTCTGCT  ACAAGTTTGT  GATAAAGGAT  TAATTTTCTT
91861    AGTTACTATA  TTTTGCAAGA  ATCAACATTA  TTATCTTTAA  ACAAAATTAA  GAATGCCTTT
91921    GTTCTCCAGA  TATAGGGATA  TCTGGACACT  CCTAAGTCTG  AGTCTGTTTA  GTAAACATTA
91981    TTTATTTGTT  CCCTTAACCG  TAAACATCTA  GAAGCTAGGA  ATGACTGACT  TTCTGGGAAT
92041    GCAGCCCAGA  AAGTCTCAGC  CTCATTTTCC  TAGCCCTCAC  TCAAAATGGA  GTTACTCTGG
92101    TTCAAGTAAC  TCTGACACTT  TTCTTCTCTT  TTTTTCTTCT  TTTTTCCTTC  CTTTATTTTT
92161    TATTTTTTAT  TTTTGAAATA  AGAAATCAAG  AATACTTGAT  GTTTCATCTA  AAACAATACC
92221    CATAATTGAT  AAGCCAAAAC  AAAAACCTAG  GTCTTCTAAC  TCAAAACTAG  GATGTTTTGC
92281    TGTCTCTGCT  GATACTCGGC  TGATCGTTAA  TAGGTAATTA  ACAAACAAGC  CTTGCTATGT
92341    CCCCCTCAGT  TTATTACCAT  TAGATCATAT  GCCTACTGTC  AATCATATTA  ATCCACAACT
92401    ATGCATTTCA  CAAAACTTGC  CATAAAAATT  CACAGGTTTC  CCGCTTCCCT  CGAGTTTTCA
92461    TTTCCGAAGG  GTCCCATGTA  ATATAAAACT  TATATTAAAT  ACATTTGTAT  GCTTTTCTCT
92521    TGCTAATCTT  TTTTTTTGTT  TTTTGAGACT  GAGCCTTGCT  CTGTCACCCA  GGCTGGAGTG
92581    CAATGGCGCG  ATCTCGGCTC  ACTGCAACCT  CCGCTTCCCA  GGTTCAAGCG  ATTCTACTGC
92641    CTCGCCCTCC  CGAGTAGCTG  GGACCACAGA  TACGTGCCAC  CATGCCCCGC  TAATTTTTGT
92701    ATTTTTAGTA  GAGACAGGGT  TTCACCGTGT  TGGCCAGGAT  GTTCTCAATC  TCCTTACCTC
92761    GTGATCCGCC  CGCCTCGTCC  TGCCAAAGTG  CTCGGATTAC  AGACGTGAGC  CACTGCACCC
92821    GACCAATCTG  TCTTTTTGTA  GAGGGGCCTC  AAGCATGAAC  TTACTGATGG  GTGAGAAAAA
92881    CAGAATTTTC  TTTTCCCCTA  CAATATAAAC  ATTAATTGTA  ATGTTATCAT  TCAGGACATT
92941    TTGGTGACCA  ATCTTACAGA  AATTTTATCT  TGTGCAAGTC  TATGCAAACC  AATATGTAAA
93001    TCTTCTATAA  GTGAGATTGT  ATTTCACTTT  TCTAGTATCC  TTTTAAATTA  ATAAAAGAGA
93061    TTCTAATGAT  TATTTTCATT  ACTGCATTTC  ATTGTAGGGA  AGTAGATAAT  TGCCCTTTAT
93121    TCACTGACCT  TCGCTTTTTA  AAAATTTAAA  CCATGTTACC  ATGAAAATGC  TTTTCAGTAT
93181    TTCTCTACAC  ACAAGATTGC  TGTAAGGGCA  AAAATAGAGA  TAGGAATCAT  GCATCCATTG
93241    ATATACATAT  TTTGATTTTT  AATACATGTT  ACCAAGTTGC  CTCCTGAAGG  TCTGTTTACA
93301    CTCTCACCAA  CAGGGTGTTT  TTTCCTGACT  TCCACAAATG  CTCTTGAACA  GTGGGTGTGT
93361    TAGTCTGTTC  AAATTGCCGA  CATGAACAAT  TAAATCTCAT  TGTTGTTTTT  ATTTTTAAGA
93421    CAATTATTGT  TTGAGACTGC  ACATTTGAT   AATAACATTT  CTTCTATTAT  GGTTTGATTA
93481    CTCATGATTC  TTGCCCATTT  TCTTTTGGGA  TGTTGCCTTA  TGTACATTAT  TTTAAATAGA
93541    TAGCTCCATG  TATTAAAAGA  TTATTAAGTT  TGAGGGCTTA  TGATATGTCA  GTTACATTTC
93601    TAAGATTTTT  TTTTTTTTTT  TTTTGAGAC   GGAGTTTCAC  ACTTGTTGCC  CAGGCTGGAG
93661    TGCAATGGTG  CGATCTCGGC  TCACCGCAAC  CTCCGCCTCC  AGGGTTCAAG  CAATTCTCCT
93721    GCCTCAGCCT  CCCCAGTAAT  TGGGACTACT  GGCAAGCGCC  ACCACGCCTG  GCTAATTTTG
93781    TATTTTTATT  AGAGATGAGG  TTTCTCCATG  TTGGTCAGAC  TGGTCTCGAA  CTGCCGACCT
93841    CAGGTGATCC  ACCCGCCTCG  GCCTCCCAAA  GTGCTGGGAT  TACAGGTATG  AGCCACTGGG
93901    CCCGGCCACA  TTTCTAAATT  CTTTATAAGT  ATAAATTCAT  TCAATCTTCA  CCAAAACTCA
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93961   ATGAAGTGTG AGTACTATTA TTATCATTGT TTTACAGATC AAAACAAGTA ATACAGTCAC
94021   TTACTGAGTT CTATACACCT GGTAATTTTT TTGTTTCGTT GTTCTATCAA TTATTGGGGA
94081   AGGGGTGTTG AAATCTCTAC CTTTAAATCA TGTATGTGTC TATTTCTCCT TTCGGTTCTA
94141   TCAGGTTTTG CTACACATAT TTTGCAGTTC TGTTATTTGG TGCATATACA TTTAGAATTG
94201   CTTGTTTTTC GTATTGGATT GACCCTGTTA TCATTATGTA ATATCCCTGT CTGTTCCTAG
94261   TAATTTTCTT TGCTCTGAAA TATACTTATC TGATATATCA TCCAAAAGAC CACCAGGATG
94321   GCTAAAGAGT AGAAAGGAGA GATTTACTGG CAATACTAAT TTGCAAGCCA GGAAGAGATG
94381   GTCCCAGAAC CTGCCAAAAT TACTCTCTCT TTGGGGAGAA GGAGCAGGTT GGTTATTTTT
94441   ATGCCTCATA GGCTATATAT TACACAATAG AGTCATACAT ATTTAGCACG TTTGGGGGGA
94501   CAGCTATATA TATTATGAGG GGTGCCAAGT GCATTCACAA TGGATAAACA CGTGTAATAT
94561   ACCTCCCATG TTCACTTCGA GGTTAAATTT TGGTTAAAAT GAGGTAGAAT TTAGGTCTTT
94621   ACATCACAAG GTGAACTATA GGAACAAAGT TTACGTGCTG CCTCTAGCAG CTGGCTGAAA
94681   ATGGCTTAAG GTCTACAATT ACGTGTAAGA ATAGAATGTG TGTCAAGGCG GTCCTCTGTC
94741   CAATCAGAGT TGTAGTGGAC TGGACTGTAA ATCAGAGTTA GGAGGGCTTC TGATAGCTCC
94801   TATAGTTAAG GAATTAGCA AGTGTGAGTT TTTTGGTAGT CTTTGGAATT TAGGAATTTG
94861   CCATGCCAGC CAAGCCATGA ATGCTCTACC AGTAGGTAAC TTTGTTTGCT TAATCTTAGA
94921   GTCTGTCTTA GTTGGTATAG GGGCATCTAT TTTGGTCTTT CAGATCCCAG ATATTATTAA
94981   TACAGATACT CTTGCAGTTT TGGGCTGATG TTTATATGGC TTATCTTTTT TGCAGCCTTT
95041   AATTTCAACC TGCGTTATGT TTATATTTGA AGTGAGATTC TTGCAGACAG TGTACAGTTG
95101   TTGTTTTTTT TTTTTGAGA TGGAATTTCA CTCTTGTTGT CCAGGCTGGG GTGCAGTGGC
95161   ACAGTCTCAG CTCACTGCAA CCTCCGCCTC CTGGGTTCAA GGGATTCTCC TGCCTCAGCC
95221   TCTTGAGCAG CTGGGATTGC AGCCATGCGC CACCACACCC GGCTAATTTT TGTATTTTTA
95281   GTAGAGACAG GATTCACCAT GTTGCCCAGG CTGGTCTCGA ACTCCTGACC TCAAGTGATC
95341   CGCCAGCCTC GGCCTACCAA AGTGCTGGGA TTACAGGTGT GAGACCTCGC GCCCAGCCAA
95401   ACTGTTTTTT TATGGGTGTA TTTATACCAC ACACATTTAA TGCAATTATT GATATCTTAG
95461   GGCTTAAGTT CATGAAGGGT AGTGTGGGAA CCATAGTCTC TTGGCCCACT AAATGTTTGC
95521   CAGAAATCAC TGACAAGGCA GATTGATTAA TAGGTGAAAA GGCATTTTAC CTATTGTTTA
95581   ACGTGTCTAT GTGGGAGCAT TCAGAATTAA TTACCTAACT TCCCAATGAG TTATAGATGC
95641   TTATATACCA TTTTTAGATC ACAGAAAGAA TTGGGGCTTA GATTCTGGTA AAACAGGTTA
95701   TGGGAGGCAA AAGAGGTTTG GCTTGCAAAG GTGGCCTTGT TAGGTAGGTG AAGCCTCCCT
95761   CAGAAAGAAC AGATGGTAAA TGTTTCTTTT ATGATTTTTA AGTGTCAGAC TCTCAGTCTC
95821   TCCTGGATCT GGGGAAAGGT ATAGAAAGGT GAGGAGGCAT GGCTGCATTA ATGGAGATTC
95881   TCTACAGATG TAAAATTTTT CCCATTTAAG GCAGCTTTGC AAGCCCATTT CTGCCTGCTG
95941   GCCAAGCAGC AGCCATTTCA AAATATGTCA AAGAAATATA TTTTGGGGTA AAATATTTTG
96001   ATTTCCTTTA GACTGGTGGC CTTATAAGAA AAGGAAGAGA CACCTGAGCT GACACACATA
96061   CCCTTGCTCT CTCAACATGT TATGATGCAG TAAGAAGGCC CTCACCAGAT ACTAATTCCA
96121   TGCCCTTAGC TTCCCAGGTT CTAGAACAGT AGGAAATAAA TTTCTTTTCT TTAAAAGTTA
96181   GCCAGTCTGT GGTATTCTGT TATAGTATCA CAAAATGGAC TAAGTAACTA TATTATGATC
96241   ATCTTACATG ACTGATCCCT CCTACATCAT ACACATACAC AGGCCACATT TGGAACATTG
96301   TTAGAGGTTC CTCTGCCCAG TACAAATGTA CTACAAATTA TATATGTATT TTTAAATTTT
96361   TGAGTATCTT CAATAGTATA TTTTCGTTAA CTTTTGTAGT CAAAATGTCA TTATAACATG
96421   TATTCAATAT GCATAATTAT TAGTCAGATG TTTTACATTC TTTCTTCATA CTAAGTGATA
96481   TGGTTTGGAT ATTTGTCCCC TCTAAATCTC ATGTTGAAAT GTAATCTCCA ATGTTGGAAG
96541   TGAAGCCTGG TGAAAGGTTT TTGGATCGTG AGGGTGAACC CCTCATGAAG CGCACTCTTC
96601   AGGGTAATCA ATGGGTTCTC ACTTTGAGTT CACAAGAGAT CTGGTTCTTT AAAAGAGTGT
96661   GACACCTCCC CCATCTCTCT CGCTCAGCTC TCACCATATG ATATGCCTAC TCCCTCTTCA
96721   CCTTCCACCA TGATTGGAAG TTTCCTGAGG ACTTGCCAGT AGCAGATGCC TGCACCACAC
96781   CTCCTGTACA GCCTGCACAA CCGTGAGCCA AAAAAAATTA CTTTTCTTTA TAAATTAGTC
96841   AGTTTCAGGG ATTCCCTTAT AGTAATGCAA GAACGAACTA ACACACTAAG TCTATTTCAT
96901   ATTTACAGAA TAGCTCAATC TGAAGTACCC TTTTTCAACT TCACAGTAGC TACTTGTAGC
96961   TAGTGGGCAC TGATTTGGAG CGTGTTCAAG GGTGAATTGT ATTATGCAAT TAACAGATTT
97021   TTTTTATTGT TTTCGCAAAC CACGAGGCAT AGATTGTCTT ACTTTCTCTG CTCCTGGTGT
97081   TGGAGTTGTT ATTGGGAAAC AACTTATTTT CCTCTTATAT TTATATGGAA TAAATAACCC
97141   CCAATATTTC CCTCCCCAAT ATCTGCCTTT TGTATGTTTT TTGAAGGCAA GTGCCTAGAA
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97201   TTTACTGTTT TTGAAGCACT TACTGAAAGG ATTGCCATCA AGTTGTTTTG CTAATAGTAC
97261   ATGCCAGGCG CTTGTTGGTT TGCTTAATTC AAGGTAACTT GGATGAGAAG AAGAGTTTTT
97321   CTCATCCATG GCTCAGTGGA GTATAGATTA CTGATATTGT GACTGGATGT ACTCCTGCTT
97381   TCTAGTCTGA GTTTTTGAAG CTACCCTTAA TCTTGGTTTC AATTTTATCT AGCCCTGTAC
97441   ATATCCAAGG CTCTTTCCAA AATGGTCTAC GATTGTTTA GGAAGTTAGA ATAGCTGTAC
97501   TTTCTGAACC ACGGTTCCTG ACATTTTCTG GACTTCAAAC ACATCCAGCA TTTTATCGAA
97561   GTATTTATCC TTCCTACTTG GCTGGCTTCT TCCTTGCCTT CAGGTCTGAA TTCAAATGAC
97621   ATTCTCCTGA TGAAACTTTC CATCCTTATT TCTATTCTTT TTTCTTATCC CCTTTCTTTA
97681   TTTTTCTCCA CAGCACTCAT CACTTATCTC TACATTTTCA TTATGTATTT ACCTTATTGT
97741   GCACCTCCCA CTACAAGACA AGTAGCACCG TAAGGAAACA GGTTGTCTGC TTTTTCACTG
97801   CTATGCTCCC TGCACCTAGA ACACTCTCTG GCACTTAGCA GGTTTTCAGT AAATATATGC
97861   TGAACTAATA ATGCTGGATA TACATCTCCC TCATGAACTC TCTAAATCCT TCTAATTTAC
97921   ATTGATCAAT CTTCTTTTCC ATGTGCTTTT GTATGATTTA TTGCTCAAAA TCTTTATTTT
97981   ATATGCAGAA CGTGCACTGC TATTTAATCT TCATGTACGT AAGTCCTCCC TTCTCTGAGT
98041   ATAATCTCTT CAGGGCACTA TCTGAGATAA CTTTTTAACA TCTCCATCAT GAATCTTGTA
98101   CCTTTTCAAA GAAAATGAGC CAGTGATTAC TGATGTTTAC GGCTATTGTT GAGGGTGAAG
98161   ATCATTATAA TTTTGAAAAG GGAAGTTGAA TATTGTGAAG GGAAAGATAA CACTAGAGTC
98221   AGAAGACTTG GGAGAAGGCA AAAAACAAAC TAAAAATGAG CACTTTTAGT CTCCTGACAG
98281   TTTCTCTGAA TCAAATCCAT AGTTCTGTGA CAGCGTTGGC TTAGAAGCAG ATTTTTTTTT
98341   TTTTTTTTTT TGAAATGGAG TTTCGCTCTT GCCCAGGCTG GAGTGCAGTG GCACGATCTC
98401   GGCTCACTGC AACCTCTGTC TCCAGGGTTC AAGCGATTCT CCTGCTTCAG CCTATGGAGT
98461   AGCTGGGATT ACAGGCTCCC ACAACCACGC CCAGCTAATT TTTTGTATTT TTAGTGAAGA
98521   CTGGGGTTTC ACCATGTTGG CCAGGCTGGT TACGAACTCC TGTTCTCAAG TGATCTGCCC
98581   GCCTTGGCCT CCCAAAGTGT TGGGATTACA GGCATCAGCC ACCGTGCCCA GCCAGGAGCA
98641   GATTTTTTTA CACTCATGTT TCTTTTTCCT TCTGTCATCC TGTTTCAGTA TAAGCAGACC
98701   ACAGATAGAA GTAGTAGATA CCTCAGAAAT TCCTGGAATA ATTAATCCAC GTTCATCTGT
98761   ACTCCATCTG CTCCTATCTC ATGGAATATA AAAGGAAAAA CACCAAGATT TCCCTAGGCA
98821   ATCTGTCTTG ATTTTAGGTT CCTCAACAGG AGAGCCAGAC AATGGCTGTA ATAATATTGT
98881   CCCGGCCAAG GAAAAACTTC CCCTTTGCCC TCCCAAGGTT TATGGAAAAT TACTGGCAAA
98941   ACACAGATTA ACTGGAGAAA AGGCATATAT ATTTATTTCA TCACAATTTT ACAGGAGATT
99001   TTAGAATTAA GACTGAAAGA TACAGGGGAA ATTGCCCATT TTTATGCTTA GGTTCAACAA
99061   GATAAACAGC TGTATAGGGT ACGATCTAAT GCTAACAGAC TGAGTGGGGA AGCCCCGCAA
99121   GGCTTGTCTG TCAAGATTCT TCTTGACCTC TCAGTGCAGC ATTTCTTCCT TCTGGTTATA
99181   GGACAAGACT CTCTTTTAGA ATGGGGGGTC TTATGACCTA CAGGCAAACA AGGTAGGTTA
99241   GAGTAATACT TTTAGGTTTT ATGGCTGGTT CTAGGGAAAA GGAGTTCTGG TTTGTATGGC
99301   CTACCTTGAG GAGGAATTCT GGTTTCTATG GCTAGACTTT GGGGAGAATG GGACTTACAG
99361   ACAGGAAGGC AGAAGGTGGT CAGTGAAACA CTTTTATAAT CATAATCCCA TTTTGAGTAT
99421   TTCTGTGTTA TGGAATGTTT GTTCTCTCAT TTCCTGAAAG ATTCCAGAGA CTCCTCATTC
99481   AGTGTTGTGA AAAAGTTCAG GAAATGCAAC TCAAAAATGT GCCACTTTGT TACGCTGATT
99541   TCTTTGAACT GAGGGCACCT AGGAAACAGT AAATTCAAGG AAGGGCTTTC GCTGAACTCT
99601   AATCAAAAAT TTGAAAATTA AAAAAAAATT CAAAAAGGAA TTTAGTTGTT AAGATTCACT
99661   TCCCTGGGGA ATCTCATCAA CCAGAGAAGA TTAACTGTAT CACAGGAGAG GAGACTGGTG
99721   GTTAACACCA TCTAAACAGA CTTTGTCACA GCTGTCACCT ATTCTTTGAA ACACCCATTT
99781   ATTTTTCTCC AAAATCATAT ACTCTCCCCT AAGTTGCCTA CATCCCCCTT CTTTCTCCCT
99841   TATGAATCAA GAGAGCTTAT AAGCTTCTAC AGTTCACTGG GATTTGGGGT ATTCGCTTTT
99901   CTTCCCTCCC ACTCCCCCTC CCCTTTTTTT GTCTTTGAGA CACAGTCTTC TGGCTCTGTC
99961   GCCCACGCTG GAGTGTGGTG GCTCTATGTG AACTCACTGC AACCTCCTCC TCTCGGGTTC
100021  AAGCGATCCT CCCACCTCAG CTTCTCGAGT AACTGGAACT ACAGGCGTGC ACTACCAAGC
100081  CCGGCTTTTT TTTTTCTTTT TCTCCCCCGT TTCTTTTTTG GTTATTTTAC TGGAGACAGG
100141  GTTTCTCCAT GTTGTCCACG CTGGTCTCGA ACGCCTGACC CGCCGTCCTC GGCCTCCCAA
100201  AGTGCTGGTA TTACGGGCAT GAGCCACTGC GCCCGATTTG AAGGACCTCT TAAATATCTA
100261  TTTAGAAATT GGTCGGAGTC CACTCCTTTC CAAAAACATG AGTCACAATC CGGGAAAAGC
100321  ACGAGCGGCT GAAAGTCAAA ATAACCAGAA CAAAACCTCC ACTCATGCTT AAAAAAGGTA
100381  TTTTGACAAA ATCCTAATTC GGCCAATTAT TATTAGTATT CAAGTCGAAG GCTCGTCAAG
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100441   CCAGACTGGG GATTGGGTCA AACATAAACC TTACACCAGA CGGAAGGATT ACATGCAAAT
100501   GAAGGATGCA GATTCTGATT TCCCATTGGG TATTTGACAT TAGCCAATGG GAGAATTCCT
100561   CACAGCCTAC CTCCAGTCAG TATAAATACT TCTCTGCCTT GCGTTCTAAT GTAGTTTCAT
100621   TACATTTTCT TGTGGCGATT TTCCCTTATC AGAAGTAGTT ATGTCTGGTC GCGGCAAACA
100681   AGGCGGTAAA GCTCGCGCCA AGGCTAAGAC TCGGTCTTCT CGTGCAGGTT TGCAGTTTCC
100741   TGTGGGCCGA GTGCACCGCC TGCTCCGCAA AGGCAACTAC TCCGAGCGCG TCGGGGCTGG
100801   CGCGCCGGTG TATCTCGCGG CGGTGCTTGA GTACCTGACC GCCGAGATCC TGGAGCTGGC
100861   GGGCAATGCG GCCCGCGACA ACAAGAAGAC CCGCATCATC CCGCGCCACC TGCAATTGGC
100921   CATCCGCAAT GACGAGGAGC TTAATAAACT CTTGGGGCGT GTGACCATCG CGCAGGGTGG
100981   CGTTTTGCCT AATATTCAGG CGGTGCTGCT GCCTAAGAAA ACTGAGAGCC ATCATAAGGC
101041   CAAGGGAAAG TGAAGAGTTA ACGCTTCATG CACTGCTGTT TTTCTGTCAG CAGACAAAAT
101101   CAGCCTAACA GCAAAGGCTC TTTTCAGAGC CACCTACGAC TTCCATTAAA TGAGCTGTTG
101161   TGCTTTGGAT TATGCCGCCC ATAAAGATGT TTTTGAGGTG TTTTTAATGG CTTTGAGTGT
101221   GGCACTTTTA GTAATTTGTC CTGCAGAAAT TAGATCCATA GAAACCTCAG GAATTCTAGG
101281   TATGTGGGAG AAGTGCCATG CAGCACAAAA CATGTTTACA GGGGTGATTC GCGTTAAGTT
101341   TCACACACAG CAGTTACTAC ATTTTAGAGG AAGGAAATTA TACCCATGAG TGCATTCCTA
101401   ACTATCTTGA ATGGAAGTGT TAAAACCCGC ATGCCCCACA CAAGTTTGAA TATGTCATAC
101461   CATTTGCTGT AGCAATTAAT GGCATACACA ATTGAGAGCA CACACATTAC CACTGAACAT
101521   TTGAGTATGT ATTTCCCAAA ATGAGCTTTT TTCCAGTTTG GGGATGTTTT GCTTTGTTTT
101581   GGGGTGGAGT CTCCCTCTCG CCCAAGCTGC AGTGCAGCGG CGTGATAACA GCTCACTGTA
101641   ACCTCGAACT CGGGCTCAAG CGATCCTCTT GACAGCCTTC TGAGTAGCTG GGATTACAGG
101701   CGAGAGCCGC CACGCCCGGC TAAGAGCATT TTTCTAATTG CCCACACTTC TTATGCGACA
101761   CCCAGAAAAA TACAATTTTA AATAAAGCGC ATATGCAAAT TTCCCTAATC GTCTCCAATA
101821   TTCTCTGATT TCTTTTTTAT ATTTTAACTA GAAACAATTG GAGGTTTCCG CGTTGCTTTG
101881   TGTGGTTGTA AATTTTAAGA CTTCAGGAAA CTTTTCCAGT ACAAGACTTG TCCACAGTGG
101941   ATATAGCAGC TAAGGGGTTA ACAAAATGAC GTCAGAGTAG CTACGGTAAT GGGCAGGAGC
102001   CTCTCTTAAT CTGCAACCAG GCACAGAGAT GGACCAATCC AAGAAGGGCG CGGGGATTTT
102061   TGAATTTTCT TGGGTCCAAT AGTTGGTGGT CTGACTCTAT AAAAGAAGAG TAGCTCTTTC
102121   CTTTCCTCCA CAGACGTCTC TGCAGGCAAG CTTTTCTGTG GTTTTGCCAT GGCTCGTACT
102181   AAACAGACAG CTCGGAAATC CACCGGCGGT AAAGCGCCAC GCAAGCAGCT GGCTACCAAG
102241   GCTGCTCGCA AGAGCGCGCC GGCTACCGGC GGCGTGAAAA AGCCTCACCG TTACCGCCCG
102301   GGCACTGTGG CTCTGCGCGA GATCCGCCGC TACCAAAAGT CGACCGAGTT GCTGATTCGG
102361   AAGCTGCCGT TCCAGCGCCT GGTGCGAGAA ATCGCCCAAG ACTTCAAGAC CGATCTTCGC
102421   TTCCAGAGCT CTGCGGTGAT GGCGCTGCAG GAGGCTTGTG AGGCCTACTT GGTAGGGCTC
102481   TTTGAGGACA CAAACCTTTG CGCCATCCAT GCTAAGCGAG TGACTATTAT GCCCAAAGAC
102541   ATCCAGCTCG CTCGCCGCAT TCGCGGAGAA AGAGCGTAAA TGTAAAGTCA CTTTTTCATC
102601   AGTCTTAAAA CCCAAAGGCT CTTTTCAGAG CCACCCACTT ATTCCAACGA AAGTAGCTGT
102661   GATAATTTTT TGTTGTCTTA ACAGAACAAA TTTCTAAGGA CCCCCCCGGA AAGCATTAGA
102721   CTATGGTCTT AAAGTTGATT AACAGAAATA ACGGTTTGGT CAGTCTTGCA GTGTAGGTTA
102781   TTTCTGACCT TATTAAGGTG CTATTGGAG AGAAGCTGTG TAAGTCCACT ATCATTCAGG
102841   CCTCTAGCTT GCTATGATTA GCATTGTTT AAACAACTTT GTAAGAGTAA GGGAAAAATC
102901   TGGTAAGTAG TTAACTGGCG CTTACTAGGC ATTTTTGCAA AGCTTTGAAA AGATTAGAAA
102961   ATTGTGTCTT GCGAGTTCCA GTGTCTTCCT CAAAATGCTT AGGAAGATTT TCTCAGCTCA
103021   ATACATAGTC CCCTAGGTTT TCTCATATAT TATATATATA TATATATATA TATATACTGT
103081   TAAATTCATT TGGCTGTTAA CATTAACCTG AAATTTATTC TGGTGCAAAA TGTGAGGCAG
103141   GGATCTAACT GGCTCTCATT TTATCCATAG CTAGCTACCC ACTTTAAATC TGTCAGTCTG
103201   TCGACCAAGC ATAATTTAAT CCCTTATATA TGAATTTTTA TATGTGTGGC TTTGCTTGTA
103261   AATAGTCTAT CTGGTTGCAT TGCTTTGTCT CCTCTAGGAC TATGCACCAT GACATGCCAC
103321   ATTCTTTTTT TCAGTACTTC TTGCCTGTAG TTATTAAAAT CTAGAATTTA CAAGTTTTAA
103381   CCATTTTCTT TCTGTTGATC TTGCTTTTCG GTTTTGGAGG TTGGGGATTG AGTACTGGAA
103441   GAAAATTTAG AGGGATGGGA ATACTGTACG CAAACAAAAG TAATATTTAC TTTAAAATTT
103501   TTATATTTTG TATTTTTTTA TCATATAGCT TTTACATCAC ATTTTACAGA CTAACTTTAG
103561   AACAACCACA GAATGTCCAA CATTAAAACT ACTAATTCCA AAGACCTTGC CTCACATTCT
103621   TTTTTACAAT AAATATTTTT TACACCTAAC ATTCTTTCTT GGCCTACATC TAGAATGTAA
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103681   ACTGATGTAC CATACTAAAA TCGCCTGACC AACTGTCAAC AACAACAAAT CACACACACA
103741   AAAGATTAAA TTTGAATTGC ATCGTTTACT TAAATTCATT TGTGTTCCAG CTTTTAATAA
103801   GGCAGTTTTT GGTTTATAAA GTAATATTTG CATTTTAAAA ATTATGAAAA TGAATATGTC
103861   AGTTTGTTTT ATGATTCGTT TTTCTTGACT CTTATACAAG CGACTCTAAC TGGCATAGAC
103921   ATTTGTTATC CACAGACAGT ATAGATATGT TAGAGATGCC AATGGACTTG GTCTATGCCA
103981   AGGTGACTAC TCACAAGCTC TGGGCCCAGC TGAAGGTCAA GTATTTTTTT TCCAGTTATA
104041   GATGTGCTGG ATCTGATGTA TAGCGCTTGA CTTTTTATAT TTTCTTTATC TGTAGGAAAC
104101   AAATGTGTTG GAGGTACTGG GTCTGACGAA TAGCATAAAA GAATAAAGTT ACATTACTGT
104161   CTGAGGATCA GATGGACAGG GGGTGGTAGC TCAGTCCAGC TATTTTCCAC TCCCTCACTT
104221   ACATTCTTTG CCCCCTCCTC AACAGAACAA GGATTCTGCT GTAACTCTTC ATTGACAGTT
104281   GATATTTAAA AATTAACGAA TGGATGAAAT TCTCATTGT GAAAGAAAAT TTATTGAGCA
104341   TTTTGTATTT GTGAGTAGTG CAAACATTTT AATATTATAT TAAGAATCTA TTGTTTTGTA
104401   TTAGAGGAGT AATTAAGGAG AGATTGGAGA CAAAAAGGGG GTGTTGTTTG CAGAATATAC
104461   CATCCAAAAA TAGACCACTG TGGGATCAGG ATTCTTTTGA GCTAAAGGCA CTTCAAAAAC
104521   AGCATTCAAG AAGGGAATTC TTCTAAACTT TTCTTTCTGA AAACAGGAGA TAAAAGTTCC
104581   AATGTGAAAA ATGCTCTGCT TGTACCAGGT GAAAAGACAT ATTCTTCAGC CCAGAGGCAT
104641   AGATGAGATA ATTCTGCACA AACACAGCAG GGAGTCATAG CCGAGAGACT TCTATACACA
104701   AACAAACCTT GTTAAAATAA TCATATATTC CTTTAATCTC CTCATATGGT TTACTTTCCC
104761   ACAATTGCCT CTCTTTAACT TAATGTGAAA GCATTAGCT TTTGCCATTT CTTTGGGGCT
104821   TCACTTTTTT ATGAGGGTTC TCCTGTCCCA TAAAATTTAC ATTAAATACA TTTGTATGCT
104881   TTCATTCTGC TAATCTGTTT TATGGCAAAT GAATTATCAG GTCCAGCTGG AGACCCTAAC
104941   AGAGTAGAGG TAAAATTTTG CCTCCCTACA AGATAGAGAT TGTGTGCATT AAATGTTGTT
105001   TGTTCCCAGT TGTTCAGTTT GTCAGGCCTC TGAGCCGAAG CTAAGCCATC ATATCCCCTG
105061   TGAACTGCAC GTATGCCTCT AGATGGCCTG AAGTAACTGA AGAAACACAA AAGAAGTGAA
105121   AATGCCCTGT TCCTGCCTTA ACTGATGACA TTACCTTGTG AAATTCCTTC TCCTGGCTCA
105181   TCCTGACTCA AAAGCTCCCC CACTGAGCAC CTTGTGACCC CCACCCCTGC CAGCCAGAGA
105241   ACAACCCCCT TTGACTGTAA TTTTCCACTA TCTACCCAAA TCTTATAAAA CGGACCCACC
105301   CCATCTCCCT TCGCTGACTC TTTTCGGACT CAGCCCGCCT GCACCCAGGT AGAATAAACA
105361   GCCTTGTTGC TCACACAAAC CCTGTTTGAT GGTCTCTTCA CACGGACGCG CCTGAAACAG
105421   TTTAACAGGG TTTTTCCTGC CCAGTCACAA CAAAGTGATG TTATGCTGCA GGCTGAAGTT
105481   TACAGCTAAT GCTGTTGAAG TCTAAAATCA GTTTTGGTTT GTTAGATTTG GGTGAGATGG
105541   CTAAGATTCT CAGAGAAAGA AGTCAAGTTT GGGGTGCATT TTTCAGACTT AAAAATTTAG
105601   CAGTAGCCCT TGCAGTTTTT CCAATAGAAG TGATTAAGA ATGTTTTCAG GAAATTTAAA
105661   ACAACAGTGA GAAGCGTGTA TGGAGAGTTG AACTACACTC CAGACTTGGC TATAGGAAAG
105721   CACGAATGCT GCTATTGTAT TGCACCTTGG AAAAGAGAAC AAAGGAATAT TTTCGGACAA
105781   TTTTAACATG TCACATATGA AAAGCTAAAC GGAATCTGTC AACACCTTGT ACGTTATTAC
105841   AGGCTGTGAT TTTAAAAAAA CAATCCTTAC TAATACATAC ATAGTTGCTG CTAGCAATAT
105901   AGTGTTGGGA GTAAAAACAC GAAAATGAGA GTTCAGGACA ATATCCCAAC TCTGAGCAGA
105961   TTTTTTTAAG TAGTAACATC TAAAATTAAA CCATATTATG TAATATTTAT TTCTTTTCCA
106021   CAGTCTCTTC TCATGCCTCG TTCACATTAG CTAATTAAAA GTCCCTGAG TATCATCATA
106081   ACCCGATTTA CAGATGAAGG CACGGTTGCA ATGAGCTATC ACCCTCTTCT GAATGAGACA
106141   GTACAGTGTG AAGGATAGCA AAACTCCACT CCCATCCTCT TAGGGCTCTG GCTGGACCAG
106201   CAAATTAAAT TAATGTAAAA TGGATTAACA GGAGAAAGGT ATATGCATTT ATTTAACACA
106261   GGTTTTACGT GACACAGGTG CTCTCATAAG GTAATGAAAG CCCAAAAAAA GCAGTTAGCT
106321   ACTTATATAA TGAATTGGAC AATTAGTAAA ATGTAAAAAT GCGCTAAAGC AAAGGGATTT
106381   AGGCTAGAAT ATATAACTGT GTAGAGAAGC GCCCAGCAAG GGCTAGTGCA AGGTTTGTAC
106441   AGAATTCTCT TGGCCTCAGC CTCCTATCCT TGAGAAGAAT GTTGCTTTTT TTAAACTACA
106501   GTGAGAACAT CTTTCATATG AGAATTTCAC CTACTGCTTC TAAGAAACAG GTCAGCTTTC
106561   AAGAAACAT AAGGCCAGAG TGATCTTTTC ACGCCTGCTC TTTTAAGTAC CTTTGAATAG
106621   TCAATATGTC TTCAAGCACT TGAAAGACTT AAAAAGTTTA CCACTCCGGC ATATTAGTGA
106681   AAGCCCTTAA TATAAGCCCT TATTAAAATT CTCAGTCGAG GGTATAAATT CAGATTCAAA
106741   TAGTAGTGTC GTAAACGGGA GGGAAAAACT AAAGGGATTA AAAAGTGAAA CTATTGTGTT
106801   CTCCCTCGCA GTCCTTAGGT CACTGCCCCT CGAGGGGCGG AGCAAAAAGT GAGGCAGCAA
106861   CGCCTCCTTA TCCTCGCTCC CGCTTTCAGT TCTCAATAAG GTCCGATGTT CGTGTATAAA
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106921   TGCTCGTGGC  TTGCTTTCTT  TTCGCGTACC  TGGTTTTTGT  TGTCAGCTGG  TTAGACATGT
106981   CTGGTCGCGG  CAAAGGCGGT  AAAGGTTTGG  GTAAGGGAGG  TGCCAAGCGT  CACCGAAAAG
107041   TGCTGCGGGA  TAACATCCAA  GGCATCACCA  AACCGGCCAT  TCGGCGCCTT  GCTAGGCGTG
107101   GTGGGGTTAA  GCGAATTTCC  GGTTTGATTT  ATGAGGAGAC  TCGTGGCGTT  CTCAAGGTGT
107161   TTCTGGAGAA  CGTGATCCGG  GACGCCGTGA  CCTACACGGA  GCACGCCAAG  CGCAAGACTG
107221   TCACTGCCAT  GGATGTGGTT  TACGCGCTCA  AGCGTCAAGG  ACGCACTCTG  TACGGCTTCG
107281   GCGGTTAATC  TTTTCGTCAG  TTTTCTTCCA  ATGGCCCTTT  TCAGGGCCGC  CCACTCCCTC
107341   TCAGAAAGAG  CTGTGATTGT  ATTCTTTCGG  ATGGTAACAT  CTCAATGGCT  TTACTCGGCT
107401   ATTCTGCCTA  GTATGTAGAA  CTATTATAAA  CCAGTTGGGA  GAGACCAGGT  TGTTTGGTCT
107461   GAGTGGCTGC  TAAAGCAGAA  ATCAGCTAAG  TAAACGAGGT  CTCCGAGATA  AGTGAGCTAT
107521   AAACTTCAAT  GCTATAGTTT  TGACATGTCA  AGCAACTTAA  CGTGCAGCGC  GAGTCCGATA
107581   AATGAGTAGC  TCAGCTTTTT  AGTTTTAAAA  ACGAGTTGTG  CGTTATTTGT  ACGAGAGCCT
107641   AAGATGCTAG  CTGCCTGGAA  CTGAGTAGGT  GGATTAAAAT  GGGTGTCAGG  TCTGTTTTCC
107701   CAGGCGTATC  TGACTTAACG  TCAGCAAAAG  CTGTACTTTT  AGCTTCCCTG  GTAACACCTG
107761   CCGTCCTTAA  CCGCCCCCTG  CCGGTAGCGC  CAGAAGCCTT  TACTTCCATT  TCTAGTTGAG
107821   CTTGGCGTCC  TGCTGAGTGA  CGTCACCTCC  CCCTTCTCTG  GAGTAGGACT  GGCGGTTAAA
107881   GCTGCTTTGC  TATTTTCAGT  CCTCAGGCTG  GAGGCTCCCC  TAAGCAGGCT  GCCTACGCAG
107941   TTCGTAAATT  CCCACTTAGT  AGACTAAGGG  AGTCTGTTTT  ATAAATAAGG  ACTCAAATTT
108001   CTTCTGACTC  CGAGGTCCGT  GGCAGCAGCT  ATAAGATGGA  AGCCCCCTCT  GATGTAAGAT
108061   TCTCAGATGA  CTTGCATCTT  CACTGTACCT  GTCAACCCAA  TAGTCTTCTA  TTCCTGCCTT
108121   AAATTGTAAA  TTCCAAAACT  GATTTAATTG  TGAAAGTTTC  AAACTGTACG  ACCTAGGAAG
108181   TGTCAAAGTT  AGGTGACCAG  ATTTTTAGAA  GTCAGCCAAA  TATTCAGCAT  CTTTGATTTA
108241   GTAACAAATA  TATTGATGGC  TACTTCAGCA  AAAAAAATCA  ACTTTGTTTT  CTGGTTACTT
108301   TGCTAACAAG  CTTCTCCTGA  CAGGAGGATA  TAGTGAATAG  GCAGTTGAAT  AAGTGAGTTC
108361   GGGTGAGAGG  TCTGAGCTGG  AGATAAAAAT  GTGTGAGTCA  TCAGCAGATA  AATAAATGCT
108421   GAGACCAGAT  GAGATGGCTA  AAAACTGAAA  CATAATGTAG  TGCAGCATTG  TTTGTAATAG
108481   TAAATGAGTG  GCAACTGTAA  AGTTTTCATC  AGAAAGGACT  AGAGTGATCT  ATACATCCAT
108541   AAAATAGAGT  ATTTCTCTAC  ACAGCCCTAC  TAAAGAATGA  GAAAGCTGTA  CTCCACTACA
108601   TACTCTGGTG  TACTCTGGCT  CAGTTCTTGG  ACTCCTCTTT  TCTTGGCTAA  CTCAACTGGC
108661   CTCACCACTT  ACATGCTCTG  TGCTCTGTCA  AATAGTTTGT  TCAACAGAAC  ACCACGGCCT
108721   AGCTGTAAGT  GCCACGTTAA  CTTCTAGCAA  TGCCAAAGCC  TGTGATAGTG  GCAGCTTCGG
108781   GCTGTTTCTC  ATTCCCGGGA  TGCCTAACCA  CCTCTCCAAA  TTCTATCAGT  TTGCTTCCAC
108841   CCACTTCAAG  CTTCAGAACG  AAACATAGAG  CTTAAGAAAT  ATAGGCCCGG  CAAGGTGGCT
108901   CACGCCTGTA  ATCCCGGCAC  TTTGGAAAGC  TGAGCCTGGT  GGATCACCTG  GGGTCAGGGG
108961   TTCGAGACCA  GCCTGGCCAA  TATTGTGAAA  CCCCGTCTCT  ACTAAAAAAA  AAAAAAAAAT
109021   TAGCTGGGCA  TGGTTGCGGG  CGACTGTAAT  CCAAGCTACT  CGGGAGGGTG  AGACAGGAGA
109081   ATAGCTTGAA  CTCGGGAGGC  AGAAGTTGCA  GTGAGTTGAG  ATCGCGCTAT  TACACTTAGG
109141   CCTGGGAGAC  AAGAGTGAAA  CTGTGTCTCT  AAATAAGTGT  TTGCAATTAT  AAACCATCTC
109201   CCTGACCTTA  AATCTCTAGA  CTCATATACA  ACTGCATATT  TGATGTATCT  AATTGAATAA
109261   TGGGCATCTC  GAACTTGTCC  AAAATATGTT  TATACGTAAA  CACCAAGTCT  GTTCTTCCTC
109321   TGATATTTGT  CATGTCAATC  AATAGAACTC  CATTCTTCAA  GCAGCTTGGG  CCAGGAATTG
109381   TGCAATATTG  TTTGTCCTGA  GCTTCTTACA  ACTTTCACCC  AATGCAGTCA  GCTCTGTTGA
109441   AAATCAATCA  GAATACCTTT  CATTGTTTTC  TTTGCTGCTT  CTCTAGGAGC  AAGCTGCCAT
109501   GGCGGTTTGT  CTGAATGACC  ACAGTGACCC  CAAACTGGTC  TTTGTTTTCA  CTTTTAATCC
109561   CCCTGTCATA  CAGTTTTTTC  TCTATCCAGC  ATCAACAGTG  ATCCTTTTTG  AAGGTATTAT
109621   GTCCACTGTC  TGCTGAAAAG  ATTCCACTGG  CTTTCCATCA  CCTTCATAAT  AAAAACCAGC
109681   ATCCTTATCA  TAGCCTACAA  GTAAGATGAC  CAACCATTAC  AGTTTGCCTG  ACTCTCAGGG
109741   GTTTCTCAGG  GTGTAAGACT  TACAGTGCTG  AAACTTAGAA  AGTTCCAAGC  AAACTAGGAT
109801   GAGCTGCTCA  ACCTACTAGA  TCTGTACTCT  GGCTACCCTC  TGACCTCATT  CTCTTCGCAG
109861   TTCTTTCTCT  TCACTGACCT  TGCTGTTTCT  GGAATGGACC  AAGCATTTCC  AGCATCAGCA
109921   CCTTTATATC  TATTCTTTCT  CCCTAGAAGG  GTCTTGTCCT  GGATATCTGA  ATGGCTCTAG
109981   ATCTCATTTC  ATTCAAGCCT  CTCCTCAAAT  ACCAACCTTA  CGAAAGAGAC  CTCCCATAAT
110041   CATCCCTTGT  AAAATAAGCT  TTTCTGCTCA  TTTAGCATAT  ATATATATAG  TTGACTATCC
110101   TCAATAGCAT  ATATATATAA  CATTTCCCCA  CCTAGAATTA  TATATGTAAT  AAATATATTTA
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110161  ACAAAAAATA CATATAACTA GATATATTTT ATTTTGTGTT TGTTCTCTCT CCCCCAACTG
110221  GAATATATTT TTTGAAGGTA GGGACTTTGT TTTGTCCCAG AAGTATCCCT AGCACCTTGA
110281  ACAGGGCTGA CGTTTAACAG GTAGTTTATG GAGGTTTGTT GAATGAAAGG ATGTGTGAAT
110341  TTTCTATGTA AGTCTCCAGG CTCTCCACTA AGCCCACCAG AATGCTAACA CAATCAATTC
110401  CCCATCTCAT TCCTTGACCT GCCACTGCCT GAAGCAATCA GCGTGCAGTT TCTCTTTAGA
110461  AAATCTGGGG GATAGTCTAG GGGTTGCAAA TTAAGCAACA TTATCTTTGT TCTGAACAAG
110521  GACTGCATGA GTGTTAGGAC TGAAGAAGGC CCAAGGTGGT GGTGGGTATG CCTAAGATGA
110581  GTATGACATA TCAGCAATGC TATGAACATA GCAATGCTAT GAAAGGCCAG GCAAAACGTA
110641  ACAGGAGCTA GTCGTGGCTT ATTGTTACAA CGACTATACC TCCCATATGG GTAATCGATA
110701  TCCACACACC CCTCTACATT GACTCTGGAA TTCAGGAAAG GGAATTAAAA TTTTCTAACT
110761  TATGTACCCC AATGATTTCA ACAATATCTG GCATATGAGA TCAATAAATA TCTTTAAAAT
110821  ACCAACTAAG AAAGACATAA AATGACCCAC CCTCCATACC AGGCTCATTT TTGCTCCTCT
110881  GATTCCTGAA ACTATCCAGA ATGCAGCTAT GAATTCTCTC CATTGTCAGT TTTAAATTAA
110941  GCCAAGCTGG GTACTTGTGT AATTCCTCAA GAAATCCTGG ATGAAAACTG TCAGGTGGAA
111001  AACAGGACCT CAAAATAAAG AGACATCCAT CACTGAAGCT AACATCGTGA GGCTGAAATC
111061  AGTCCTATAA CAATGGTACC AAAAAGAGCA CAATGAGAGG CATTTGTGAA TATTTACTCA
111121  GATGAGAGTA AGATATTTCC CTATCAGCTA ACCTGAAGTT CACATCCCTT TTCCAGCTGA
111181  GTTCTGAAGC TAGATGTACT TAACTGGAAC ACATAACTGC ATCAGGAACA TCCTTTAAAA
111241  CTATGGCTAC CATGGCTTGA CTGGACAAAC CCCAGGCTTC CAGGTTAGC ACAGGTGGCC
111301  CTTCACAGAC CAACATTGCC TATGCTACCA ACCTCATGTC CTACCACCCT GCTTGCATCA
111361  TTTCTCTCTC TGCATATATA AAAATATATG TGTATGTATA TAATCAGCTT TATTGATATT
111421  TAATGTACCA CAAAATTTGC CCACTTTAGG TACAGTTCAA TGAATTTTAC CGTGTTTTCT
111481  TAGTTGTACA ACCATCATCA CAATTTAATT TCGGAATATT TCTATCACCC AAATTTCCAT
111541  TTCTGCGTAA AGGGGGAAAA AAAAGGTTA ACTGCTGAAG GCCGCGGTAA CACTGAAAAA
111601  GGTGCCTTTT CTCTCTAAAA CAGATTTTAA TCTCCCCTGA ATTTAGTGTC CTGGGTATTC
111661  CAGGAGTCTG AATAGGGTTT CAATTTTCAG GGTCTTTTTA ATAGAGTAAA ACTGTATTGG
111721  TGGCGATAAA TTTAGTATTG CTCTCAGTAC ATGATTGAGG GATACTTAAA TGTCTCTGTG
111781  ATTTTATTTC ATAATCGCTA AAAGATGGTT TTTTTTTTTC CTAAAACAGG GTTTTTGTTT
111841  TTTCTCAATA AGCTTCTTAG CTTCCCCTCC GGCTCCCTGG CTTGCCTCAG GAAATATTAG
111901  CTCATCAGTT CTGATTGGTT GACAGCTACG AATGGCCCTC ATTGATTGGG CAGCGCTTCT
111961  TTGTCCCTTG GAAACTAATA CAAATTTTTA ACACTACTTT TTTTCCACTC TTTCTTCAGA
112021  GTTGGAATAT CGTTGCTCCC CTACCCATAT GTAGTGAGTG GAGGGCAAAC TTGGAGTTCC
112081  CCTAATCTTT CCTTTTTAGG ATGTCAGCTC AGTATCATTC ATCTTAATTA CACATTGAGC
112141  TTCTTGACTT AATGGATACA GCTCTTCTTT TGTTTAGTTG GGCGGCCCTG AAAAGGGCCT
112201  TTGGTTCAGA AATGCAAGCT GTGGAGAAAT CAGCAACCTT AACCGCCAAA GCCATAAAGG
112261  GTGCGTCCCT GGCGCTTAAG CGCGTAGACC ACGTCCATGG CAGTGACTGT CTTGCGCTTG
112321  GCGTGCTCCG TATAGGTGAC AGCGTCACGG ATCACGTTCT CCAAAAACAC CTTGAGCACC
112381  CCGCGAGTCT CCTCGTAGAT CAGACCAGAG ATCCGCTTCA CACCGCCACG CCGGGCCAGA
112441  CGCCGGATGG CCGGCTTGGT GATGCCCTGG ATGTTGTCAC GCAACACCTT GCGGTGGCGC
112501  TTGGCACCCC CCTTACCCAA ACCCTTCCCG CCCTTACCAC GTCCAGACAT GACTTCCCAA
112561  GAAGTGAACC AAGAGCAAGT GAGAGAATAG GAAACCGATC TTTATATATC TACGTTACCC
112621  CTGCCCCCAC CTCCAGCGGA CACTGAGACT GAAAAGCGCG CAGGCGGGAA ATGTGACGCC
112681  TACAGTCCGC TCCTTTAACC CCTCCTCCAA GCCCCAGGAA ATGGCGGGAG CAGCGATTGG
112741  GGGAGGGTGG GGAGATGAGG GTGGGACCAA GCAGGCTTGA CCAATGGCCT TTATTTTCTT
112801  AACAGAGCTA CAGGCTTTGA GGAACTGGGT TAAGAATTAA ATGTAAACCC ATTCTGACTC
112861  CAGAATTATT TTAAGTCGAA CTTTTTTTTT AACCGAATCT CTCTGTCGCC CAGACTGGAG
112921  TACATTAGAG CCATCTCGAT TCACTGAAAC CTCTGCCTCT CAGGTTCAAG TGTTTCTCCT
112981  GCCTCAGCCT TCAGAGTGTA GCTGGGATTA CAAGCGCTCG CCGTCGCGCC CGGCGTGTTT
113041  TTGTATTTTT CGTAGAGACG GGATTCGGCC ATGTTGGCCA GGCTGATCCC GAACTCCTGA
113101  TTTCTGGTAA TCCGCCCGCC TCAGCCTCTC AAAGTGCTTG AATTACAGGC GTGAGTCACC
113161  GCGACCGGCC GAAATCGATT GGTTTTGAAG CCTTCAGTAG CATTAAAACG AAAAGTGCTC
113221  CCAATGCATT CCCTTTTGTC TTAAATTGGT TTCTTACAGC TACTTTACTT GAAAAGGTGG
113281  TGGCTCTGAA AAGAGCCTTT GCTTGGACCG TCAGAGAGAC CACAGTAATC ACGCCCTCTC
113341  TCCGCGGATG CGGCGGGCGA GCTGGATGTC CTTGGGCATG ATAGTGACGC GCTTGGCGTG
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113401  GATGGCGCAC  AGGTTAGTGT  CCTCAAATAG  CCCTACCAAG  TAGGCCTCGC  ACGCCTCCTG
113461  CAGAGCCATC  ACAGCGGAGC  TCTGGAAACG  CAGGTCTGTT  TTAAAGTCCT  GCGCAATCTC
113521  GCGCACCAGG  CGCTGGAAAG  GTAGTTTACG  AATAAGCAGT  TCAGTGGACT  TCTGATAACG
113581  GCGGATCTCG  CGCAGAGCCA  CGGTGCCCGG  CCGGTAGCGG  TGGGGCTTTT  TCACGCCGCC
113641  GGTGGCCGGA  GCGCTTTTGC  GGGCTGCCTT  AGTGGCCAAC  TGTTTGCGTG  GCGCCTTGCC
113701  ACCAGTAGAC  TTCCGAGCAG  TTTGCTTAGT  GCGAGCCATG  ACGGAAAAAC  AGCACAGCGG
113761  AACACCCAAC  ACTAGCGCAA  ATACGCCCAT  GAGCTGCTCT  ATTTATAGTG  TGTAAAGTGC
113821  AGTGATTGGA  TGATAGAAGA  CGCTAAATAT  GACGTTACAC  ACTCTGATTG  GTCTATCTTT
113881  AAGCCAGCAA  CAATCGTGCA  GTTTCACCGG  CTACTATATT  CTATTCCAAC  TCTACAGATG
113941  ATTATTTAAG  TGGTATTTTA  TTACTACTAT  TATTTTATTT  TACTTTTGCT  TTGTTCCCCA
114001  AGCTGGTCTT  AAACTTGGGC  TCAAAGGATC  TTCCCGCCTC  AGCATCCAGA  GTAGCTGGGA
114061  TTACAGGGGA  GCCCCACTGC  GCCGGCTTGG  ACTTTAATTT  TTTAAACTTG  TCCTCTTCTA
114121  CATCTGGTTT  TCATAACCTG  AAGGCTGTGT  TTATTTTCCA  TAAAACAAGG  CATTGATTCC
114181  AAAGGTATTA  TAATTCCCCA  ATTCCGTATA  ACCTTCAGCT  CTTTAGGAAA  AAAAAAAAAA
114241  AAAAAAAAAA  GAGGGAATAC  TGCTCACCTC  CTCTCCGGAA  ATGTACCCTT  TACGGGAATT
114301  TCTGAAACCT  TTCACAAGAA  TTGGATTCCT  TTGTAATGCT  TTAATTGACT  TAGGAGTGTT
114361  ATTGAAATCT  ACAAAGCATC  TCAAACATAG  TAGGATTACA  CTATTACTCA  GAAACATTTT
114421  CTATGAGACG  TCTTTCTCTT  GATTATGCTC  TTTGAATCCT  AAACTTGCAG  CGTTCTGCAG
114481  CTTTTGTTTT  CTAAAGCCTA  GGTGTACTCT  GCCAGTCACA  AAATGGCGTT  TCTCCAGCAC
114541  TGCCGCCAGG  TACCACCAGC  TGGGAGTTGT  TCCTCTTGCG  GAGCAGGAGG  TGGACTTGGC
114601  CCAAGAGAAA  CTGGATAGTG  GTTCGCAAGG  AACATAATTT  AGCATTGCCA  AGAGCTAATG
114661  CAATCATTTT  GAAAATCTCA  AAACACTGAA  AAGTGGATTG  TGACCTTTTT  AAATTCACAA
114721  GAGACAGGCC  ACATTCTATC  TTTTGATTGG  TTTAGGCTAT  TTTCTTGAAC  AGCCATTTAG
114781  AAAGCAGATC  TATCATCCTT  CATTTGCATG  GAGCGTTCCC  ATTTTATTTG  AAACCAGTTT
114841  AACCCAATAG  AAAAAAGGGA  GGCAGAACCC  ATTATTTAAA  GTGGAAACTC  CTGAATCAGA
114901  TAATTAGGAG  TATTTCCTTT  TCAAAAGTTG  CGTTTTTTCA  GATACCTCGC  TTATTACACT
114961  AAGAAAGGTT  TATATCTTTC  ACAAAGGGTT  TACTTACAAA  AATCTTCCAA  TTTTGTATAC
115021  CTGTGTTTCA  TAACTGACTA  GCCGTCAAAC  CAAGATGTAG  AGTTTCCAAC  CGTTATTTTC
115081  CAAATTTTTA  GAAATTACGT  GAAATATTTG  AATGCATGCC  TTCTCAATAA  AATGGGACGT
115141  AGGAAGCACT  GGTGCAGAAG  ATGGGTACAA  TACTTATCTG  GGACCACTCC  ATTATTTGGT
115201  TGGCACGTTG  TTTGAACAAA  AAGGGGAAAA  GCTCAGGTTA  CTTAGCATGG  TTCGGACTTA
115261  TTTGAAAACT  ACCACAGCAG  GAGCGGAAAT  AAGACCGCAT  TACCTCACTC  TCTGCTGTGC
115321  TGTGCTAGGG  GGTTATCCAG  AATAGGATTG  TAGAAGTGGA  TGTCGATTTA  ATAGTTTTTT
115381  ATTCTCCCAT  TAGCTGAGTC  TCTGATTGGC  AATGTGAGAT  CGTTTTAGCT  TATTGATACT
115441  TTGAAATGCA  CTTAACAGCC  ACAAACAAGT  TAAAGGGTTG  TTACCATAAA  ATCTTATCCC
115501  CAGGGTGTGC  TTGCATTTAT  CACCCGTGTT  TGCTTTCACA  CTAAGTGGAC  TTAACTCCCC
115561  AGCAGAATGC  CTGTCAGGGA  ACCGGTTTCG  TGGACCCAGC  ATTTAACGCC  TTTCGCAGGC
115621  TTGTGAGGCC  CATAAATATT  TGTTGAATAA  AAGAATGAGT  TGACCATGTC  ATGGTGCGCT
115681  GATTGCGTGT  GCTGACATGG  AACACAGGTT  GTAAACCTTA  ATACCAATTT  GGGGCATGTT
115741  GTATGGATGA  AAAGGGCATT  GGAAATTCCT  GAAGTGCATC  CCACATTGGA  CTGTGGAAAT
115801  AAGTTGCAAG  TGCAGAAACG  TTTCCACACT  TGCAGTTTGA  GTATTAATTG  CAGCGTTTGT
115861  GAATTCTGGT  GTTGTCTACG  ATTCATTCTT  GTTTGACGTG  AAAGGTATTC  GCGAGACACA
115921  TCGCTCTAAA  ACATTGCCAG  AAAATGTAAT  AGAGTTGATG  ACAACTGGCC  CTAACACGGC
115981  CTAAAACTCG  CACTTTTCTC  TCCCTCCGCA  ACTATTCAAA  ACACTGTATT  TTACATTTCT
116041  TGCAAATTAA  AAACTAACAT  CTCTGGCAAC  GGACCTCTAA  AAATTTCTAA  TAAAACTCCT
116101  CGGATGCTTG  TGGCACTGCA  TTTGTAAACC  GCCCCCTCTC  AACCTACTCC  CTAAAAAAGA
116161  GCTGCTTTTT  GAGAGAGAAG  CGGTACCCTC  TGATGTTACT  GGGCGGCAGT  CTGCCTACAA
116221  TTTCCTTCAC  AATGAGGCAA  CCAGAGCGGC  TTTTTCTGTG  TGTTTGCTTG  CGTTGAGGGG
116281  AGCAGGACCA  TAGGCCCTAG  AGGCCCCCAG  CTGCCTTCTG  AGACTGGGCG  AAACCCTCGG
116341  CAGCGCGCAG  GGGGCGCTAG  GGCGCGAGGG  GCGGGCACTG  ACGGGCACCA  ATCACGGCGC
116401  AGTCCCACCC  TATAAATAGG  CTGCGTTGGG  GCCTTTTTTT  CGCATCCTGC  TTCGTCAGGT
116461  TTATACCACT  TTATTTGGTG  TGCTGTGTTA  GTCACCATGT  CTGAAACAGT  GCCTCCCGCC
116521  CCCGCCGCTT  CTGCTGCTCC  TGAGAAACCT  TTAGCTGGCA  AGAAGGCAAA  GAAACCTGCT
116581  AAGGCTGCAG  CAGCCTCCAA  GAAAAAACCC  GCTGGCCCTT  CCGTGTCAGA  GCTGATCGTG
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116641  CAGGCTGCTT  CCTCCTCTAA  GGAGCGTGGT  GGTGTGTCGT  TGGCAGCTCT  TAAAAAGGCG
116701  CTGGCGGCCG  CAGGCTACGA  CGTGGAGAAG  AACAACAGCC  GCATTAAGCT  GGGCATTAAG
116761  AGCCTGGTAA  GCAAGGGAAC  GTTGGTGCAG  ACAAAGGGTA  CCGGAGCCTC  GGGTTCCTTC
116821  AAGCTCAACA  AGAAGGCGTC  CTCCGTGGAA  ACCAAGCCCG  GCGCCTCAAA  GGTGGCTACA
116881  AAAACTAAGG  CAACGGGTGC  ATCTAAAAAG  CTCAAAAAGG  CCACGGGGGC  TAGCAAAAAG
116941  AGCGTCAAGA  CTCCGAAAAA  GGCTAAAAAG  CCTGCGGCAA  CAAGGAAATC  CTCCAAGAAT
117001  CCAAAAAAAC  CCAAAACTGT  AAAGCCCAAG  AAAGTAGCTA  AAAGCCCTGC  TAAAGCTAAG
117061  GCTGTAAAAC  CCAAGGCGGC  CAAGGCTAGG  GTGACGAAGC  CAAAGACTGC  CAAACCCAAG
117121  AAAGCGGCAC  CCAAGAAAAA  GTAAATTCAG  TTAGAAGTTT  CTTCTAGTAA  CCCAACGGCT
117181  CTTTTAAGAG  CCACCTACGC  ATTTCAGGAA  AAGAGCTGTA  GTACACAGAT  GAAATCCCCC
117241  AAGCAAATGC  AACACGCCCT  CAATTATATT  AGAATCACTT  GGAGAGTCGA  TAGAACTTTA
117301  ACATAGCCTC  ATCTAGTAAG  AATTTACTAC  TCAATCTATC  AAAGATAGCA  AGGTGAATTC
117361  AAATGCACCG  AGTTAAAATC  GAGTTTTAAA  GTCACCTGGG  TTTCGGTAGC  CGGAAGTCCC
117421  GCGTCTCACG  ACTCCAAGCT  AATTAGTCAT  AACCGTATTG  AACCAAGGTT  GAAGCCCAGT
117481  CCCAGGCTTG  AGGCTTTTTA  TTATACAAGG  TTAAAGTGGG  GATATTGCGT  TTTGGGGTCA
117541  ATATTGCTAA  AGTAGCATTT  TCCGAAATTG  GGTGGTCCTA  AGAAATGCTT  CTGGGATAGT
117601  TGGCAAAATA  TATGGCTTAA  CCACGCCCTC  TCCACAGGAG  TGGCTAGCGA  GCTGTCTGTC
117661  CTTGGGAAGG  ACGGTGACCC  TGCTGGCGTG  GCTGGCGCCC  ACGTTGGCGT  CCTCTGAAAG
117721  CCCCGCCAGG  TAGGCCTAGC  TCGCTTGCTT  TCTGCAGCGC  CATCATGACA  AAGCTTTGAA
117781  ACGCAAAATG  CTTTCTTTGT  GCAGCGCCTT  ACCATGGGTG  CACTTACGGG  CTGTCGACTT
117841  GGTTTAGGCC  CTTGTCAGGA  CAAAGGAGCT  TAGTTTGTTG  GAGTTTTAGA  GCTGCAACCC
117901  AAAATCCCTT  GCTCGGTTTC  TCTGTTTTTA  GAAACGGAAG  CGCCCTGATT  GGATATTTGA
117961  AAATTACTGT  GCTTAACTGG  ATCGTGTTTC  ATCAGTCGTG  CAGGATTTTC  AACCCTGGTG
118021  GAGCCCACAC  ATTCAAAACT  GAAGATCCTT  TTCTCAGAAC  TGCCCCTTTA  AGCTTTTGCA
118081  ATTTTAATTC  TGGGGGTCAG  ATTTTAATAA  TTGGACTTTT  TTGTTACAT   CTGACAAGAG
118141  TATATGATGA  GCCAAGTTTA  CTCACTTTTA  CTTAGTGCAG  TTCAATTCTA  AAAGTTTATT
118201  TTTGCGTGTG  TGCATATGAG  TTAATAATCA  GTTGTATTTT  TCAAACGGTC  TTTTTTCAAT
118261  TGTTTTGCTT  AGCTCCTTCC  ATCGTCTAAA  GTCAGGGATA  CAGGCACATC  ACATCCCTGT
118321  TCCCCCTTCC  TCAAACTAAT  ATGTAGCTAC  CTAGGTTTAT  CCTTTAAAAC  AAAAATTCTC
118381  ACCTATTTTT  GTGAGAAATA  TACATGTTTT  TCTTTGAACT  AAGTATTTTA  CATACACCTA
118441  TCTATATACA  TGCATACTTG  TGGTTTTGTT  TTTTTAAAAA  AAAAAAAAAA  AAAACACGTT
118501  ATCTTTTGAG  ACTGGGTCTC  AGTCTGTTGC  CCAGACTGGA  CTGCAGTGGC  ATAATCACAG
118561  CACACTGTAA  CCTCCAACTC  CTGGGCTCAG  GCTATCCTGC  AGCCTCAGCA  TCCGGAGTAG
118621  CTGGGATTGC  ATGCACGCAC  CACCAAGCCG  GGCTTTTTGT  TTTTATTTTT  TGTGGAGACA
118681  GTCACACCAT  GTTGTCCAAG  CTGGTCTAGA  AATGGCCTCA  AGTGATCATC  GACCTCCCAA
118741  AGTGTTGGGA  TTACGGTCAC  TGTGCCTGGC  CTTGTATGCA  TAATTGTTTT  GTCTTTTGAT
118801  TAGGGTTATT  AATTTAAAAA  ACAAAGCCTG  GACGCAGTGG  CTCACATCTG  TAATCCCAGC
118861  ACTTTAGGAA  GCCAGATGGG  CAGATTACTT  GAGCTCAGGA  GTTCAAGACC  AGCCTGGGCA
118921  ACATGGTGAA  ATCCCATCTT  GACAAAAAAT  ACAAAAAATT  AGCAAGGCCC  AGTGGCACGC
118981  ACTTATAGTC  CCAGCTACTT  GGGAGGCTGG  GGTGGGAAGA  TGACTGGAAC  CTGGGAGGTA
119041  GAGGCTGCAG  TGAGCAGAGA  TCGTGCCACT  GCACTCAAGC  CTAGGTGACA  GAATGAGACC
119101  CAGTCTCAAA  ACAAAAATAA  TAAAAATTTT  TTACAACGAT  GTTATATACA  CTTCTGCATG
119161  TTGCTTTTCT  CTTAACCAAA  CTTTTCTAAA  ACCTGTCAT   GAAAAAGAA   ATCCTTCACA
119221  TGGAATAGCA  TAAGTTATTC  ATCCATTTCT  TATTGATAAG  CATTGATGTT  CCAGTTACC
119281  ACTGCTGAAC  ATGGTGCAAT  TGAATAGAAT  TCCAGGGCTG  AGATTGCTAG  GTTTTAGGTT
119341  GTATTTATT   ATTTTATTTA  TTTATTTATT  TATTTAGACA  GAGTCTTACT  CTGTCACCCA
119401  TGGTGGAGTA  CAGTGCCATG  ACCTCAGTTG  CAACCTTTGC  CTCCTGAGTT  CAAGCGATTC
119461  TCATGCCTCT  GGTCTCCCGA  GTAGCTGGGA  TTACAGGCAC  CTGCCACCAG  GCCTGGCTAA
119521  TTTTTGTATT  TTTAGGAGAG  ATGGGGTTTC  ACCATGTTGG  CCAGACTGGT  CTCAAACTCC
119581  TGGCCTCAAG  TGATCTGGCC  ACCTCGGCCT  CCCGAAGTGC  TGGGATTACA  GGTGTGAGCC
119641  ATGGCGCCAG  ACCTGGACTT  TGTCTTCTGT  TTCATCAGTC  CTTCTGTTGG  TTCAAGCACA
119701  GTATCACACT  GAAGACTGAT  GATTCTATAT  AAATATGGTA  AAGACTGTAC  ACCCTAACTG
119761  TTCTTATTTT  TTAATTTTAA  GGCAATTTTA  GATTCCAGCT  TTCCAAAGAA  TTGTGGAATG
119821  CTTAGAGCTA  GAGAAGCCTT  GGAAGTCATT  TAGTTTTTGT  TTTGTCAGAG  AAAATTCTGT
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119881   AGAGACTCTG  TCCTGCTCTC  ACTGAATACC  ATCCCATAGT  ACCCCCCAAC  AGCTTTAAAG
119941   GGCAATAATA  CCTTATGGAC  AGTATGCTTT  TCCTCAAATA  TATTCTAAGC  CATGGTCAAT
120001   GCAAAAGAGT  GAGAAGGAAA  GTAGAATAAG  TTATCTAAGA  ATCAGTGGGT  GCTCTCTTTA
120061   AACTGATTTA  TCACTCCCCC  TTCCAAACTC  TCTTGAAGGT  CACTCTGCCT  CCCTTTCTAC
120121   ATAAGAACTC  CTAACTCCAA  GGGAGGAAGG  TAAGTTATTC  TTATTCCTTG  CTTAGAAAAA
120181   GAGAAAATAG  GTTTGGTAAG  CATCCGCTTT  CTGCTACCAT  TCTCTGTGTT  TCTGTGTTTT
120241   TTATAGGATC  ATTCAATTAT  TGGTTGGCTC  TTGAGAGGGA  ATGCAAGGTT  CAAGGACACA
120301   AGCCTAGATC  TTGCCTGTAT  AGAACCTCAT  GATGTTATGC  TTCTCTAAAA  TGAGGCCTGG
120361   AGGAGACATG  TTGAAAGTGA  CCCATAAATC  TGCAGTATCT  CATGTCTCTC  AATGGGGACA
120421   AGGAGTACCA  TGGGAAATAG  CATTAGGTCA  ATGACAGTAA  CAACTCCCAG  GTGAGTTGAT
120481   TTATTCTTTT  ATTTATAAAG  TTGTTAATAT  GCTACATAGT  CCCTAATTTT  GCCACAAATA
120541   GTCATTATTT  TAATTTCATA  TTTCACTATT  GATAAATGAA  GGAAAAAATG  AGTAGCAGTT
120601   AAGCAGTCCA  TAAACCTACA  TATAAAGCAA  ATTGGAGATT  TTAAAATTGA  TTCTGGATGC
120661   TTAAAATCCT  TCTCATTGAA  AAAAAATTTC  GTATTAGAAG  ATTTCAACAT  TCTTTAAACT
120721   GAGAAGCATA  ACATATAAAC  AGAAAACCAC  AGCAAAACAA  AAATGCAAAG  CTCAATAAAT
120781   GAACACAAAG  TGAACACCAT  AATAATTGCC  ACACAAGTAA  AAAAACAGAA  AATCAGCCAA
120841   CCCTCCCAGA  GCCGCCTGAT  GCTTGCTTCC  AGTCACATTA  TCACTCCATC  TGCCCTAAAC
120901   ATAACCCCTA  TTTTGATTTC  CAATGCTGTA  ATTTAGTATG  CCTGTTTTTG  AAACATATAA
120961   AATGGAAATA  AAACAAATGT  AATCCTATGT  ACCTGACATA  TTTCACTCCA  GAACATTAGG
121021   TTTGAATAGA  TTCATCTGTG  TTGCTGTGTA  TAACTTTAAT  TCATTTTTAT  TGTTATGTAA
121081   TATTCCATGT  TATGAGTGCA  ACAATTTAGG  TGTCTACTGT  TGATGCATAT  TTGCTTCCCT
121141   TTTTCAGCTA  ATATAAACAA  TACCGTGAAT  ATTCCTGTGT  ATGTGTCTTG  GTATATATAG
121201   GAATACATAT  TTTGTTTGTA  TACCTAGGAG  AGGAATTGTT  GGGTCAAATG  CTAAACTCTT
121261   TTTGAAAGTG  GTGATATTAG  GTTTACATGC  GATGAAATGA  AAATTAAAAC  CACAGTTATA
121321   AACAGCATGG  ATGAACCTCA  CAAACCTAAT  GTTGATGGAA  TCTAGCTGGG  AATTCCTGTT
121381   CTTCCATATA  CTTCCCAATA  TTTTTTTCCA  ATTAAAATTG  TTAATCTTTT  GAAGATGTTA
121441   TCCATTGTGG  CAGATGTGCA  GTATTATCTC  ATTATGGTTT  TATTTTACAT  CTTTTGCCCA
121501   TTTTTTCTTA  ATTGGATTGT  ATATCAGTCG  ACTTGGGCTG  CCATAACAAA  AATACTAGAC
121561   TAGGTAGCTT  GAACAAAAGG  AGTTTATTAC  CTCACAGTTC  TAAAGGCCAG  GCCAGAAATC
121621   CTAAATTGAG  GTGCCAAGAG  ATTCAGTTTC  TAGTGAGGGC  TCTCTTATTG  ACCTGAAGAT
121681   AGTTGCTGTC  TTAGATTGTT  TGGTGCTGAA  CAGAATACCA  GAGACCAAAT  AATTTATAAA
121741   GAATACAGAT  TTATTTCTTA  CAATTCTGGT  GGCTATAAAG  CCTATGGTCG  AGGGGCCCAC
121801   CTCTGGCAAG  GGCCTTCTTA  CTGTTATGGC  AGATGTGAGA  TGTCATCTCA  TATTCAAACC
121861   ACAGCAGTCG  CCTTTTGTGT  CCTCATGTGG  CCTCTTCATA  TGCCCATAAA  ATGACCTCAT
121921   GTCTCTTCCT  TTTCTTATAA  GGACACCAGA  TCTATCAGAC  TACTGGCCTA  CTCTTATGAC
121981   CTCATTTAAC  CTTAAATATC  TCCATAAAGT  CCCAAAATCC  CTATCTCCAA  ATATAGGCAC
122041   ATTGGGTGTT  AGAGTTTCAA  CATCAATTTT  GGGGGAACAC  AATTTAGGCC  AAAAAGATTG
122101   TGTTTTTTCT  TGTTGGTTTA  AGATAGCTGT  CTTTTTGTCC  TTTTTGTCCT  TTCTTTTTTT
122161   TTGAGGTGGA  CTCTTGCTGT  GTCACCCGGG  TTGGAGTGCA  GTGGCGCTGT  CTCAGCTCAC
122221   TGCAACCTCC  ACCTCCTGGG  TTCAAGAAAT  TCTCCTCCTC  CCAAGTAGCT  GGGACTACAG
122281   GTGCATACCA  CCGCGCCCTG  CTAATTTTTG  TATTTTTGAT  AGAGACGGGG  TTTCACCATG
122341   TTGGCCAGGC  TGGTCTCAAA  CTCCTGACCT  CAGGTGATCC  ACCTGCCTCG  GCCTCCCAAA
122401   ATGCTGAGAT  TACAGGTGTG  AGCCACCAAA  CCTGGCCTGT  CTTTTCTGTT  TTAAGTTTTT
122461   AAATTTTGCT  CACGAACCCT  TTATCCATTT  TATGTGTTGC  AGGTATTTCC  TCTGTAACTT
122521   GTCTTCACTC  TGTCAGAGGC  TGGAGTGCAG  TGGCACAATC  ACAGCTCACT  GCAGCCTCCA
122581   CCTCCCAGGA  TCAAGCGATC  CTCCCATCTT  ATCCTCCTTA  GTAGGTGGGA  CTACATGTGC
122641   AGGCCACCAT  GCCCAGCTAA  TCTTTGTATT  TTTTTGTAGA  GATGGTGCTG  TTGCCCAAGT
122701   TGGTCTCAAA  CTCCTGAGCT  CAAGCAATCC  ATCAACCTTG  GCCTCCCAAA  GTGTTGGGAC
122761   TAGAGGTGTG  AGCCACCACT  GCACCAGCC  AATGATATCT  CATGATGCAT  TAAAGTCATT
122821   AATTTAGTGT  ACTCAAATTA  AGCACACTGC  CCTTTTATGC  ACAACCTTTT  TTGTATCTTA
122881   TTTAAAAAAT  CATTTCTAT   TTCAAGGTCA  TGAAGATCTT  ATTTTATAAT  ACCTTCTTGT
122941   GAAATTAGTT  CTCAAGACTA  CCCTCACTTC  TAACACCAAT  TATAAGTTGG  GAGGTCTGTG
123001   GTTCCCAATC  AACCTTAGGT  TAGTAATTTG  CTAAAAGGAC  TCACAGAACT  TGCTGAAGCT
123061   GTTAGCCTCA  TGGTTACAAT  TTATTATAGG  ATATATAGCT  TATTATGTCA  TTCCAATGCA
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123121   ATGTAAAATT ATACAACTAC TTTTAAAAAG ATTTTAGCAT TTGACCCAAC AATTTCACTC
123181   TGAGGTATAC AAACAGCAGA TATGTGTGCA CATATATACC AAGACACATA CACAGCAAAA
123241   TTCATTGTTT GTAATAGTTG AAAAGGGGAA ACAACTCAAG GAATAAAGAT TAAAATCAGC
123301   TGAGAAAAGA AACACACAAG GCAGTATTAT GGATCGAATT GTATGCAGAT CTCCCTTGCC
123361   CCCAGAAGAT ATGTTTAAAG TCCCAACTCC CAGTACCTCA GAATTGTGGC CTTATTTGGA
123421   AATAGGATAG TTGCAGATAT AATTAGTTAA GATGAGGTTA TAGTACAGTA TGATGGGCTG
123481   GTGACTTAGA AGAAGTAGTA TATATATATT TTTTAATAGA ACTAGTATTC TTCTAAGGTG
123541   GTCACGTGAA GACAGACACA CACAGGCAGA GACTGAGGTT ATGCAGCTGC AGGTCAAGGA
123601   ATGTCAAAGG TTGCCAGCAA GTACGAGAAG CTAGGAAGAG TCAAGGAAGG ATTTTCCTAC
123661   AGGCTTCAGT GGAAGCATAG ATCTAATGAT ACCTTCATGT CAGATTTCTA GCTTCCAGAA
123721   CTACAAGAGA ATATATTTGT TGTTTAAGC  CACCCTAGCT TCTAGCTCTT TGTTACAGCA
123781   GCCCTAGGAA ACTAATATAG GCACAATCCA GGCAAGTTCC AAATATGAGC TTCCAGTTGT
123841   CCTCTCCCAG TAATATGAAC AGTATTACTT TCCCAGCATT AATGTGTGAC AATACACATG
123901   ACGTACAGAG CAGTCCCCAC TTATGCACAA AACATATGTT CCAGGACCTC CAGTGGATGT
123961   CTGAAACCAT GGATAGTACT GAACTCTATA TAGCTGTTTT TTCCTATACA GACACAGCTA
124021   TGATAAGGCT TAATTTATAA ATTAGGCACA GTAAGAGATT AATAACAATA AATTAGAATA
124081   ATTGTTAAGA ATATACTGTA TAAAAGTTAG GTGAATGTTT ATTTCTGAAA TTTACCGTTT
124141   ATTATTTTTG GACTGCAGTA GACCACAGGA ACTAAAACCA TGTAGAAACC GTATACAAGA
124201   GAACTGTATT TCACCCGAGC CTCAGTGTGC AGTTTTAATG GCCTGCCATG GTTGACTGCT
124261   CACATGGCCG ATCTTTTAGT CTACCTCCAC AGGTAGAGCT GATACTGTGT GGCTCAAAGT
124321   TCCTATTATA AATCACATTG TTGACTGTGT GGTGGTCAAA ACCTCCAGGT AAACAAAGAC
124381   ACACTTATCA GTGAGAACAT TTCAAGGGTC TAAAATTCAT CTCCCAGTAG CTGAGGGCAA
124441   AGGCTAGACC TCTTTTTGGG TAAGATAAAT TTTTTACCAT ATACTTTATT TTGCTTTTCA
124501   TGTTTAACTT TATTTTGCTT TTCATGTTAG TTCCCCTGGA ATTGTTTTTT GTGTATAGTG
124561   TGAAGTAGGG GGTCAAGTTT CTTTTTTTTT CCTTTTTGTT CTTTTTCTGT TTAAAAGGCT
124621   ATACAATTGT CCCATGCCAT TTATTTACAA GAGTCCTTTC ACCATTGTTG TATGGTGCCA
124681   CTTTAGATGT AAATCAATGT CCATATTTGT TTGAGCCTGT TCCATTCGTT TGTCTATTTT
124741   TGGACAACAC TGCCCTGATT ATTGTCATTT TATCAGTTTT GATATTTAAT AAAGCAACAG
124801   ATTTGTTTAT TTTGGGCCCT TGGATTTGTG TATTAAATTT GAACCCTGTT TGTCAATTTC
124861   TATAATAAAG CTTATTGGGA ATCTGATTAG GATTACAATG GTTTTGTAGA TCAGTTTGGG
124921   GACAATTAAT ACCTTTAAAA TATTGACCGC TTCAACTGTA AATATACTCC TCCATTATTT
124981   AGTTTTCCTG TTTAATTTAT CTGAGTAATA CATTATAGTT TTCTTCGTAG AAGTCAGATA
125041   CGTAGAAAAT TCAAAGCCCA AGTGCAATAG CTCATGTCTG TAATACCAGC ACTTTGGGAG
125101   GCCGATGTGG GTGGATCACC TGAGGTCAGG AGTTTGAGAC CAGACTGGCC AACATGGTGA
125161   AACCTCATCT CTAGTAAAAA TACAAAAATT AGCTGGGTGT GGTGGCGGGC ACCTGTAATC
125221   CCAGCTAATC AGGAGACTGA GGCAGGAGAA TCGCTTGAAC CCAGGAGGCA GAGGTTGCAG
125281   TGAGCCAAGT TCCTGTCACT GCACCCCACC CTGGGCGACA GAGCGAGACT TCGTCTCAAA
125341   AAAACAAAAA AAAGAACATT CAAATAATCA ATGTAGATAA TTCAAATAAC TAAAAAATGA
125401   ACAGTTATTA AAATATCAGG ATATAAAAGC AAAAAAATCA ATAACCTCCA TATATACAAA
125461   ATGGCCAGTT AGAGAAAAAA AAAAGAATAG GCGAGACTTA AAAAGGCTGG GAATCTCCCT
125521   GAAAATCTTT GAGAGCCTTG GCCCTGCCCT CAGGGATTTC TCTGGCTTCA TGCCCAGATA
125581   CGGGTACAGT TCCTTGTTTA AAAAAATTTT GCTCCATCAA TCAACAAGGG GCTCCTTCCT
125641   CAGAGCACAA GGACCTCCAT AACACCGGAC ACTAGATGTC TAAGGGACAC CTCTTAAGGA
125701   AGTTAGACTT CCAAAGAATG GTGTTTCCTC TGTCCCCAAA CTCTGGAACT CACAGCACAA
125761   CTGCTCCTTG GAGTTCGGTT TCAAATCTAC AAGGCTGTCA TGGAGGTTGC AGACCAAGTC
125821   CGTGGCCTCA GTGTCCGGAT GTACGGTGGC CTTGGCACCT GAATGTGAGA ACATGACCTC
125881   CCTGAAACCA CCACAAGTAT TGTTTCATGT TATGTATGTT TTTTCTTATC TGAAATTCCT
125941   TTTCTTTAAA AATTCAAATT ACATATTTTG CAAGCCCCTG AACAAGCTTC ATGAGCATTT
126001   ATTGAACCCA CAGCTTTTAA AACCTACTGA ACACTTTGCT CTATGTTGTC ATTCACTATC
126061   CACCAATTAT TTAATTATTG ATCAATATTG TTTCCTTAGT GTTGGGATCA TTTATGCATG
126121   TATTTCTTTT ATATTGCATA TTTTATATTT CTGCATTACA GTTATTACAT ATTACTTTTG
126181   CTACAGTAAT AGTTCAAAAG TGTACATCCA AAATTTAGCT GTGAAGTGGA TGGACTGAGG
126241   CAGAACTGGA GGCAAGAAAA TGTCACAGTA ATTCTAAAAA AGATGATGTA CAATTAGAGC
126301   AAGAGAGTAG CACTGAAATT GAAGAAAAAT AGATGCGTTT GAGAGAAAAT TAGGAGGTAG
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126361    AATCAACAGA TTAGATGTAG GGATGAGAAG GGTCAAAGAT GACACTAGGG TTTTTAACTG
126421    GAGCAAGTAG GTAGACAGAA CATTTCTTCC TGAAAGGGCA GGTCAGATCA TGTGTTGTCT
126481    CAAAGGGCAT GAAGAGTAGA AAGCCTGGGA CAGATCCTGA GATGACCAAT ACCCATGGTG
126541    CAGGGAGAGG GAGGGAGATC TGCTAAAAAG ACTGCAAATG TCAGGATAGT AGAAAATCAT
126601    GAGTGTGTGA TGTCCTGGAA GTTGAGACAG TATCACATTT GAGAACATTT AAATTGGTAA
126661    CTCTGACAAA AAGCTGGAGG CCAACTGTGA ATGCCCATGA GAGTGAGAAG CTCCCACACT
126721    TTTGTGGGCA TCAGAAAGCC CACCAGGTTC CTGCAGTGAA GATCTGAGAA GGATCCTCTT
126781    GTGGCTTTGG CAGGGAGAGA AGAATTATTA TGAAATACAC CCCAGAACCT TCTTCAAAAC
126841    AAAGGCCTAC TCTCAAGGGG AAAACATTTT GCCAGAGTCT TATCCCAGCT GGGAGAAGGT
126901    AATTCTTCCC ACTGCAGCCT CATCTAGGCT TTCTGTCTCA CTTAAGGGAA GAAAATTAGT
126961    CAACAGGGAT CAGAGCTTCA TGAAAATAAA TTGGAAATGG TGCAGCCAGG AAAGGAGCAA
127021    AGGTCTGAGG AGGAGGAGAA GGAGGAAGAG GAGTTGTATC ATTATAAATA CTTGAGGAAG
127081    AGGAGGAGAA GGAGGAGGAG GAGGAGTTGT ATCATTATAA ACACTTGAGG AAGAGGAGGA
127141    GGAGAAGGAG GAGGAGGAGT TGTATCATTA TAAACACTTG AGGAAGAGGA GGAGGAGAAG
127201    GAGGAGGAGG AGGAGTTGTA TCATTATAAA CACTTGTGAC GGTCCCAGCC CCAAGATATA
127261    GGCATGCTAA TAAACTGAGG CTTAACACTT TGACTACAGA ATGCTGCTTC TCCCTAACAC
127321    CATCAAGGCT CCAACTGAAT AACAATGAAT TATGAATGAA AGAGCTGTAA GGAGAGACAA
127381    AAGTTAGAAT GAGACAAGTA TTGTTATCTA GAGATGCCAA GAAGGCAAGG AAGATAACTA
127441    AAAAGGCACT CTGGATTTAG AAATAGGAAG TCATTAGTGA CCTTGTAAAT AATGGAGCCA
127501    GAGGAATACC AAGGGCAGAA GCCTCACTAT AGTGTGTTGC ACCTGTCAGA GGTCAGGAGG
127561    TGTAACTGAC TCTCCCACAG TGTGGCTTTG GAAGAGAGAA GTCAGCAGCT GCATGGAGAT
127621    TTGGGAGAGG GAAAGCTTTT TTTTTTTTT TTTAATTGGA AAAGACTGAG CTATGTGTAA
127681    ATAGAATAAG ACAGGAAGAG TGTAGACACA GGAAAGAGGG CAGACAAAAA CAAGTGCACA
127741    GTTATCTAAG GGAAACAATG GGATCAAGCT GCAAGTATAT AAACTTGTCT TGATAGAAGA
127801    ATCCTTGATC TGGTTTATTC AGTGTTTGGT CCAAACCCAC ATCCCTGTTC TGCCTGTCTC
127861    TGACTTGCTC TGTGCCCCAG AAGCCCAGCT TCTACAGATA GCATTAGCTG GGCAGCCCTG
127921    CCCTCTTGCA ACAGCTGGAT TTGGCCAGTG ATCAGCCCAG CAGGAATGTA GATGGCAAAG
127981    GAGAGAGAGG TTAGTGTACT TATTCCCTGC ATCACCCCCC TGCTTGGTGG GCAGCTCTTC
128041    CTCCACAGTC CCAGCTCTGG CCTAGCTCTG GTTACAGGTT CCCTCCCATT GCCTCTTCAG
128101    ATTTAAAGGT GTGTCTGTCA GGGTATAACT GGGAGCTAGA AATTGCACTG AAATTGAACA
128161    AAGAATTTTA TGGGAATGGT TGTTAACTAG TTATAAGAGG ACTGAAAATG GAAAAGTGGA
128221    CAAACGTATC AGAGATAGTA ATGACAGAAA GCAACTACCA CCTCCAGGTT TAGGAGAACA
128281    AGGAAAAGAT TCTTTGAAGA GATCCCCAGA ACTGGGACCT CTGAGGAGTG TATGCTGGAC
128341    CACTGATGAT GATATGTCTG TAGATAGAGG CATGATGAGG CTGATTTTAG GAGCATGGAA
128401    GATCTCCAAA CTGAAGCCAA CTGCTGTTAC TGGATTCAAC TGCCACTGCC AGGTTGAAGA
128461    ACCCATTCTG TGAGGATGTC AACAAACAAA GTGGGAAATC TTTTCACATC CTTCCAGCCC
128521    TCTAGTCTTC CTCCAGTGCT TTCTATTGGT AGGGTTTGGG GAGGTGGCTA GCAAAGCGGT
128581    ATTGGAAAAG ATAGAAGAGA CTAAATCTTC ATAACCAGCA CAGGGTGACA CTGGATCACT
128641    ACTGTTGCTG ATCTTGGGCT GCCTCATATC CCCTGTTCTT CCCATTAGCC CTGTCACAAC
128701    TTTGTAGATA TCCCTTCATT ATATGCCCTT CATATATTCT TTTGGTTTAA CTTTTTCTGT
128761    TGGAATCCTA ATATGGCACT CCTCCATTTT TCAGGACCAA AAGAGTATAA AAGATTATCT
128821    TTTACCAAAA AAAGACAAA AAACTGATCT AATTCCTGAT TTGATCATTA CACAATCTAT
128881    ACATGTATCA AAATATCACA TAGTACCCCA TAAATATATA CAACTGTGTC CATTAAAAAT
128941    AAAAATTAAA GAAAAGATGG TAAATATAGC TCTGTCAGGC AGTGGAGGTT TTACCACGAT
129001    GGCTGTTATT TCCCCCATGA AGGGGGGAGT GAGGGAGCAG CTGAAAGTAG GTGCTTATAG
129061    GGGTATAGAG GGGCTCAAAG CTTTGAGAGA GGAGAATGTC TGAAAGAGCT GCCAAATAGC
129121    ATGCAGGTCC CATGGGGGCA GAGCCTCTGC TCATTCACCA GTGCCTCTTC AATATCTACA
129181    CTTAAGCCTA ACACAAAGTG TGTGCTTAAT AAGTATTTGC TGAGTATGTA AAGTGGAAAC
129241    AGAACCAATC TGGCAAACTT TGTAGGACTG GTGGGCAATG AAGATCAGTC AGGTAAAATC
129301    TGTGGATATA AATTTATATT GATCAAAAAA TTCAAGGTTA GGTGTTTTTC TTCAGTCATG
129361    CTCAACGATG CTTCAGCCAT GCTCAACTCT TCTGTAGCCA CAGAAAAAAG TTTACCCATA
129421    ATCGAGCTGT GTCTGTGTCT GAATAATGAA AAGACCATGA TGCAAGGGAG TTGGAGACAC
129481    AGAAACAGTG TTTGAAGTAA TGGGTAATGG AAGCATGCTA CCAGGGAAAG GAAAGAAGTG
129541    GCAATAGGAA GGAACAGAGA TCTGTGGTCC TATGTCCCCT GAGCATATTC ACATGTTAAA
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129601  GCTAATTCAG TTTTCAATCA TCATTAAAAT TTTGTTCCTA AATATATGGC CATTATTTTC
129661  CACAACCACA CTAAAACTTT ATTACCTCTG GCAAGTGACT ATGCAAGTAA CTAAGAGCAA
129721  AAATATCCAC AACTACCATT TGAGCTATCA ATTTAGGGAA AGTCATCTGG CTATAATCTA
129781  AGTGACCCTC CACTGAATGT CAGTATCTTT GCATATGTGA TTTAAATCTG GGCCTTCGCA
129841  ACACCATGAA CTGTTCTTGT CTTGAATATC CAGATTGAAG GAAATAATCT GAGTAGTTAC
129901  GAGTCCTGAA GCTAGAAAGA TGGAAACCCC ATTTGCTCAT CAGAAAGCCT TAGAGCTTGG
129961  GCGCTGGCGG GTCCTGTCTC ACCGGGACAG AGGGGCTCTT TCCTCCCCAT CTGATAGTCT
130021  GATAACTAGA GAAGCCGGCC AACTTATTCT CCAAGAAGGA GCCATCTTAG TTCCTCCTGA
130081  AATGTTCATA TTTAGAAATT ATTGTTTGTC AGTAATTTAA CCCCTTAATG GGCTTGCCTT
130141  GTGGTCCATA CCACTGAGTG CAGAGCTTGC CTGGAAGAAT TGTGAGGGCC ATTCCATCTT
130201  CCAGGCAGTA GAGTTCAGTA CTTCTTTAAA ATTGCTGCTG AACTCTGTAT TTGAAAAGAA
130261  AGAATCATTT GGGTGTGGTA GCTCACACCT GTAATCCTAG CGCTTTGGGA GGCTGAGGTG
130321  GGAGGATCAT TTGATGCCAG GAGGACCACT TGAGACCACC CTGGGTAACA TAGCAAGACC
130381  CTGTCTTTAG AAAAAAAAAA TACAATAAAA TAAATACAAT AAAAATAAAA GCAAAAGAA
130441  AGAGTCCATC TTAGGGACAG ACTGTAACTA CTCACTGGAG CTTACCTTTA CATAGTTCAG
130501  GATCAATTAT AATAAACAC TTTTGTGCAG ATTCAATAGG ATTATTTTAA TCCCCATCAT
130561  CTCTCTGAGT TTCCAGTCAG TTTCTCTGCA TGTAGACACC CTTCTCCAGC CCACCATTGT
130621  CTCTCCTCCT ATAGCTCCAC CAACAAATCA GAACTTTTTC TAACTGCACC TAGTGCACCT
130681  AGAGTCTACT CCAGAATGCT CATGGAGAAA GTTTCTGAAA GGTAAAACTC TGAATGATAT
130741  TTGTAGCTAA AGGGAGACTT GCTAGAGACA ATAAGCTAAT AGTTGTAGAC TTCAGTAGAA
130801  GAGGAATGAC ACTGCAATGT CAGGGTGCAG GACTTCAAGA GGGCAGAGTA TGGAAACCCA
130861  ATGGGAAAAA TGCTCACCAG GAACATGAAG AGAAGGAATT ACGTGTAAGG ATTTCTCAAT
130921  GTGTTCCCAA ATTTGCCCAG CAGAGGGAGG CCTCGGGTTG ATGGCAGGCT GACCACACAA
130981  TTAAAGAAGG CTGAACCTGG GGGCTTTTAA CAACCATCGT GGGCTCTACT GTAAGCATTT
131041  AGAAAAAGAA AGTTATCCAT TCAAAAATAT ATATATTTTT AAACTTCAGA ACAAAATTAT
131101  GAAGAGCTAT ATTTACTTTT CTACATTCTA ATTTTTATAA ATCTGAGTAT ATTTTGCATA
131161  TATTGTTATA GTACATATTC AATTTTGTAT TTTGCTGTTT TCACTTAACC ATTTTTACTA
131221  GATTACTCTG TGTTCATAAT AATCACTTTT TTAAAACTTT TATTTTTATT TATTTATTTT
131281  TTTTTTGAGT CAGAGTCACA CTCTGTCGCC CAGGCTGGAG TGCAGTGGCG TGATCTTGGC
131341  TTACTGCAAC TTCCACCTCC TGGATTCAAG CAGTTCTCCT GCCTTAGCCT CCTGAGCAGC
131401  TGGGATTACA GGTGTGCACC ACCAAGCCCG GCTAATTTTT GTATTTTTAG TAAAGACGGG
131461  GTTTCACCAT GTTGGTCAGG CTGGTCTCCA ACTCCTGACC TCATGATCTG CCCACCTTGG
131521  CCTCCCAAAG TGCTGGGATA ATCACTTTTT ATGCTGCATA ATTCTTCAGA TTTGTCAGTA
131581  CGACTGTATT TACACTCATT TGTTTTATTA GAAAGAATTC CAGAATATTT TGGCTGCCCT
131641  AATTAATTTT ACAATTAATA TGATTTTGAA ATTGGGTATT GGCTCCTTCT GAATTGGTTT
131701  ATTAAAATAT ATTCTAATGT AATTTATGAC ATTTTCATCA TATTAGCATA TTTATTCTGT
131761  TAGAATTTCA TAATTTATAA AGCTACAAAC TGTATGTGAT ATAGCTTGTA ACTTTATCTC
131821  ATAACTTTAT GCAGTTACAA GTAGAAATAA AATGTTCCCC TCAAGATTGC TTAAAATTTT
131881  ATTATAAACA AGTGTAAAAA ACAAAATCAC TAAAACACTC CCTCTTTTTT CCCCCAAAAT
131941  GCATGTTTCC ATTTTAACAG AACCCGTATT TAATCAGCAG ATTTCTATGG TGGCTAGATT
132001  TGTAGACTAA ATATTAAAAG TCCCAAAGCA AATGCATTTT TCTCTTAAAT TTTACTGACT
132061  TTTTTTTTTT TTCTTTTTCT GAGACGGAGT CTTGCTCTGT CGCCCAGGCT GGAATGCAGT
132121  GGCACAATCT CGGCTCACTG CAACCTCCGC CTCCCGGATT CACGCCATTC TCCTGCCTCA
132181  ACCTCCCGAG TAGCTGGGAC CACAGGCGCC CGCCACCACG CCCAGCTAAT TTTTTGTATT
132241  TTTAGTAGAG ACAGGGTTTC ACCGTGTTAG CCGGGATGGT CTCGATCTCC TGACCTCATG
132301  ATCTGCCCAC CTCAGCCTCC CAAAGTGCTA GGATCACAGG CATGAGCCAC CGCGCCCCGC
132361  CTACTGACTT TTATCCAAAG AAAATATAAG AGCTCTTCAT CATAACGTAT GTTTCTTGCT
132421  CTTGTTATTA AATATGACAC ATTTAGACTT AAACTGATTT GAAGGTTTAT GACATTGTTT
132481  AAGTTATTAC ATAATTAATT CATAAAGATA ATGACTAGTT TGAACTACTG ACAGCTCACA
132541  CATCATCAGT TGAACAGCAG AAAGCTTACT AAGCTACTTT CTTATGTTTC TGTCTCCCAG
132601  CTACTAAAAG AAACGAAACC CTTCCAGGTG TTAAGGCAAA ACTTTCCTCC CCCTTTCTTC
132661  TATAAATCTG ATTCCATGTT AGTGAAATTT CTACTGATGG CTTTGGTTTC CTCTATAGTA
132721  GAATAGAGAT CCTATGGCAA AAGTCATGTC TGACATGGTA GCAAATAGAA ATGGGGAAAA
132781  GGAAGGTCTG CAAGAGCCAA TGTGGGAAAT GGGGAGAGGA CTGACTACAA AAACCCAGCA
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132841  GGAATTCCAG  AAGAAAACTC  CTCAGGACGG  GCACATTGGC  TCATGCCTGT  AATCCCAGTA
132901  CTTTGGGAGG  CCGAGGTGGG  CAGATCACTT  GAGTCCAGGA  GTTTGAGACC  AGCCTGGTCA
132961  ACATGGCGAA  ACCTCATCTC  TACAAAAAAT  AAAAAAATTT  GTCAGGCGTG  GTGGCATGCA
133021  CCTGTAGTCC  CAGCTACTCA  AGAGACTTAA  GTGGGAGAAT  CACTCGAGCC  TTGGAGGTGG
133081  AGGTTGGTGA  GCCGAGATCA  CGCCACTGCA  TTCCAGCCTG  GGCGACAAAG  TGAGACGCCA
133141  TCTCAATCAA  TCAGTCTCCT  CGAAAAGCAA  CATTATGGAG  AGACAGGATT  CCGTCAAGGC
133201  CTGGGGCACA  CAGGAAAATA  TTAAGGCAGA  AGAGAGTTTC  CTCCCCACAC  CACACCGTAT
133261  CCCACAGGCA  CTGCGGATGT  GCATATGCAA  GAGGGGTTGA  TCCTAAGAAT  TTAGAGTCAC
133321  AGAGGAGGAG  GCACCAAGCA  GACTGTGGAG  AAAGTCATGA  CCAGAAAGGG  ACAGAATGTA
133381  AAGCTTCAGC  TGATTATCTG  GCCTCAGGGA  TTCCAGAGGA  ACTGGTCCCA  ATGGTCTCCT
133441  GGTGATGTAG  GTTCTTAGGT  TTCTTTTACA  GGGGTTTTCT  GGGAGATCGT  TGACCCAGTT
133501  AGCATTCAAG  CAACTTCCAC  CCTGCACTTT  TATTCTTTCC  CCTTCACCTG  CTTAGGTTTT
133561  ATCTGTCCAG  GAAATAATAA  TAAAATTATT  GAGCCCTGGA  CATGTACCTG  TAAAGCTCCT
133621  TAAAGATGAT  GCCTTCTAAC  TCCTCATTCA  ACAGATACAA  AAACATTACA  ATAAAATGAC
133681  TCATGCAAGA  CACCCAGGTA  GTTTATAGCA  GCTAATAAAA  ACAGAATAAC  TATAAAATAT
133741  GGTAAGTTTA  TAAAAGTTAC  ATTGAGTATA  CTTTATAAGA  ACTGCTTATT  GAGTTTGCCT
133801  AATAACCACA  CAGCACAATA  ATAATATGTA  TATATTTTTA  AATATGTGTA  AATATGTGTA
133861  ACACAAACTT  GTAGAAGGTA  TATCTGAGTA  CAACCCTATT  CTGTTTGGTT  ACCTTTTCTA
133921  GTTCATTATG  TAAGTGGCAT  AGCTACCTAA  GGACTTATGC  TTATAAATGT  TACTCAAAAA
133981  AATACAGAGG  ACATATGTGG  ATAGATAATG  GAAGAGATAA  GATAGGTAGG  TTGAAGGGTT
134041  GGGCTGCCCC  TCCACACCTG  TGGTTGTTTC  TCGTTAGGTG  GAATGAGAGA  CTTGGAAAAG
134101  AAAGAGACAC  AGAGACAAAG  TATAGAGAAA  GAAAAAAAGG  GGTCCAGGGG  ACCGGTGTTC
134161  AGCATACGGA  GGATCCCACC  GGCCTCTGAG  TTCCCTTAGT  ATTTATTGAT  CATTATTGGG
134221  TGTTTCTCGG  AGAGGGGGAT  GTGGCAGGGT  CAAAGGATAA  TAGTGGAGAG  AAGGTCAGCA
134281  GGTAAACACG  TGAACAAAGG  TCTCTGCATC  ATAAACAAGG  TAAAGAATTA  AGTGCTGTGC
134341  TTTAGATATG  CATACACATA  AACATCTCAA  TGACTTGAAG  AGCAGTATTG  CTGCCAGCAT
134401  GTCCCACCTC  CAGCCCTAAG  GCAGTTTTCC  CCTATCTCAG  TAGATGGAAT  ATACAATCGG
134461  GTTTTACACT  GAGACATTCC  ATTGCCCAGG  GACGAGCAGG  AGACAGATGC  CTTCCTCTTG
134521  TCTCAACTGC  AAAGAGGCGT  TCCTTCCTCT  TTTACTAATC  CTCCTCAGCA  CAGACCCTTT
134581  ACGGGTGTCG  GGCTGGGGGA  CGGTCAGGTC  TTTCCCTTCC  CACGAGGCCA  CATTTCAGAC
134641  TATCACATGG  GGAGAAACCT  TGGACAATAC  CTGGCTTTCC  TAGGCAGAGG  TCCCTGTGGC
134701  CTTCCTCAGT  GTTTTGTGTC  CCTGAGTACT  TGAGATTAGG  GAGTGGAGAT  GACTCTTAAC
134761  GAGCATGCTG  CCTTCAAGCA  TTTCTTTAAC  AAAGCACATC  TTGCACAGCC  CTTAATCCAT
134821  TTAACCCTGA  GTTGACACAG  CATATGTCTC  AGGGAGCACA  GGGTTGGGGC  TAGGGTTAGA
134881  TTAACAGCAT  CTCAAGGCAG  AAGAATTTTT  CTTAGTACAG  AACAAAATGG  AGTCTCCTAT
134941  GTCTACTTCT  TTCTACACAG  ACACAGTAAC  AATGTGATCT  CTCTCTCTTT  TCCCCACAGG
135001  AGGTGATGGC  CGGAAGAACA  TGGCAGAGGG  CAAAACAAAA  CAGCATTGGG  AACAAGCTCT
135061  GTTTAAAAGG  AGACTTGTGA  ACAGCAAAGA  GTAGAAAGGG  TTCTCTTACA  ACTGAAGCCC
135121  ATGGAAGACA  AATGTGTACT  GCGTGAGTTT  TAAGGCAATA  GGAGTAGTGG  GACCTAGGGC
135181  ACACCAGAGA  GCATATTAAC  TCTCAAACTT  TTAAAAACAT  TATATCTGCT  GGACACAGTG
135241  GCTCACACCT  TAATCCTACA  ACTTTGGGAG  GCCGAGGCGG  GCGGGTGTAG  CTTGAGCCCA
135301  GGAGTTCGAG  ACCAACCTGG  GCAACATGGC  AAAATCCCGT  CCCTACAAAA  CAAACAAACA
135361  AAAAACAAAA  TTAGCCAGGC  ACGGTGATGC  GTACCTGTGG  TCCCAGCTAC  TCAGAGGCTG
135421  AGGTGGGAGG  ATCGCTTGAG  CCCCGGGAGG  TTAAGGCTGC  AGTGAGCCAT  GATAATGCCA
135481  CTGCATCTCA  GCCTGGGCAA  CAGAGGGAGA  ACCTGTCTCA  AAACAAAAAC  AAAAACACAC
135541  CATACCCAAC  CACAATGCAT  CTGTCTTAAG  TACCAGTACC  ACACCCCTCT  ACTCACTACT
135601  AAATAGGTGA  GTTCCCAATC  CCTGGTAGCA  GGTTTAAGCA  TGTTATATTA  AAGGTCTTAG
135661  GCTAGTGACT  CATTCACTCA  TTAAACAAAT  ACTTATTGTG  CATCTACTAT  AAACTAAGTA
135721  CTGTGCTAGG  TACAAAAGCA  AATAATCTAA  GCTCTATAAA  CTTTACTTTC  TTCATCAACA
135781  AAATGGAGAT  GTTTTAGGCA  TCTACTCATC  ATTCTGAGCT  CCATCTTTTG  TGACTGTAGT
135841  TGGCAGAGCT  TTTTATCAGT  TTCTCTAAAT  AGCTCTACCA  GTCCCTGGTG  GATGCTGGCA
135901  TGCCCAAAGG  ATCCATCCTG  ATGGCCCTGT  CTGCTTACCT  TACCTGCCTG  CCTTTGCAGC
135961  ACCGCTCTGC  TCTTCTGCAG  GACTTCCCTT  ATCCTTTGGG  GTCTTGCTGC  TCTTAGGCTG
136021  CTCTGCTTGT  TTTGATCTGC  TTTGCATCAC  ATGTATGTAA  AGGTCCTTTC  CTTATTTACC
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136081   CATGACCAAG GTATTATGAG ATTCTGGAAT TTCCCCAAAC CACATTGATT GCTGGGAGAA
136141   TAGAAGAAGT GGATTACAAG TGGAACTTAG AAGGGGAGTA TTCGAGAAGA CGTCTCTGCA
136201   AATCCATTTA GAGAGACCTT TCTCCAGTGG TGACTCAAAG ATGCAGCTCC TTTCATCCTG
136261   TGGCTTGGCC ATCTTCAGCA CATGGCTCCC AAGGATGTCC TCAGGATGGT CTCTAATCCA
136321   AGGAGCCTGA AGAGAAAAAA AGGCATGGAG TATTGTGAGT GGTAGGTGGT TATGGACCAG
136381   TTATGGAAGA ATACACATCA CTTTTGCCCA CCTTCTACTA ACCAGAACTC ACACAGCCAT
136441   AGACACTGAC AAGTAGGACT TAACAAGAAT CTAATTTTGA GTCTAGGAAT ACGACTGTAG
136501   CAAATATTTA ACAGCTTCAA ACACAGGTGC ATTGCTATCA CTATGCTTGG CCCAGGCCTG
136561   TCTCCCTTTC CTGCCATGTC ACAGGGGCCA GCATTTATGT CTAGATTGGG TTGGTTGGGA
136621   TATTAAGACA ATAATGAACC AATACAACAT CTTGAGCATA AAACCAACTG ATACAATGAT
136681   GTACAAGTCA GATGATTCTG ATGATTATGA ATTATGTCAA TAAAAGAAAT GTGATAACTA
136741   AGGTAATTTT TGTTTTGGCA AATTTTTGTT TGTTCATGAC AGGATGAAAT CCTGTCATTT
136801   GTAGCAACAT GGATGGAATT GCAGGATACT ACATTAAGTG AAATAAGCCA GAAACAGAAA
136861   GTTAAACACC ACATGTTCTC ACTTATATGC AGAAGCTAGC TAACTAAGTA AATAAGTTTA
136921   TCTCATTGAA GTAAAAGTA CAACAGAGAT TACTAGAGGC TGGGAATGGT AGGGGAAAGA
136981   GATGATAAAG AGAGATTCGT TAAAATAAGT TACAGCTAGA TAAGAGCAAT CAGTTCTAGT
137041   GTTCTATTTG TACTACAGAA TGGCAATAGT TAACAGTAAT AAATAATTTC AAAGAGCTAG
137101   AAAAGAGGAC ATTGAATGTT TCCAACACAA AGAAATGAGA AATGCTTGAA ATAATGGATA
137161   TTCTAATTAA TTACCCTGAT CTGATCACTA TACACAGTAT GTATAAAAAT AACACTATGG
137221   GCTGGGCGCA GTGGCTCACA CCTGTAATCC CAGCACTTTG GGAGGCCAAG GTAAGCAGAT
137281   CACTTGAGGT CAGGAGTTAG AGACCAGTCT GGCCAACATA GTGAAACTCC ATCCCTACTA
137341   AAAATACAAA AATCAGCCAG GCGTGGTGGC ATGTGCCTGT AATCCCAGCT ACTCAGGAGG
137401   CTGAGGCAAG AGAATTGCTT GAACCCAGGA GGCGGAGGTT GCAGTGAGCC GAAATCGCGC
137461   CACTGCACTC CAGCCTGGGT AACAGAGCAA GGCTCTGTTT CAAAAATAAA TAAATACATA
137521   AATAAATATT TTTTAAAAAA AGAACATCAC TATGCACCCC ATATATACAT ATAATTATTA
137581   TGTCAATTTG AAACATAATT TTGAAAAATG AAAAAATGAA ACACAAATAT GAATCAATCC
137641   TCTCCAAGTT GATATACTTA AAAGGAAAAA AGTCCGAGGG CTTAAACTAT TCAATCAAAA
137701   TTTTATTAAA ATGCTATAGT AATCTGGAAA GTATTTCAGA ATGAATTGGT ATAAGGTTAG
137761   ACACAAAGAT CAGTGAAACA AAACAGAGAA CCCAGAAATA GATTCACACA TCTATGGACA
137821   ACTGGTTTTG ACAAAGGTGT CAAGGCTATT TAATAAGTAA AAAAATCGTC TTTTCAGTAA
137881   ATGTTTCTTG AACAAGTAGA CATCCGGTGT GGGGGAGAGG AGCAGGAGCC TTACCTCAAA
137941   CTTTATGCAA AAATTAACTC AAAATAGACC ATAGACTTAA ATGTAAAAGC TAAAATTATA
138001   AAACTTCTTT AAAAAAATAGG AGAAAATCAT CAACACCCTA GGATTAGCAA AGATTTCTTT
138061   AAAACAAAAC AACAGGTTTA TAGTTTATAA AACATAAATA ACAAAATGAT AAATTTCATC
138121   AAAAGTGAAA ATTTGCTTTT CAAAAAACAT TATAAATGA AAAGCAGGAG GCTGAGGCAT
138181   GAGAATCACT GGAACCCGGG AGCTACAGGT TGCAGTGAGC CAAGATGGTG CCACTGCACT
138241   CCAGCCTGGG TGACAAAGTG AGACTCTTCC TAAAAAATAA ATAAATAAAT AAATAAATAG
138301   AAAAGAAAAA GAAAAATCAC AGGCTGAGAG AAAATATTTA TAATACATGT ATCTGACAAA
138361   GGACTCGCAC CTGGAAAATA TAAGGAACCT TATAACTTAG TAAGATGACA AGCCAAAACA
138421   AAGAGTAAAA GTTTTCAACA GACATTTCAC AAAAGAAAAC ATACAAATGG CCAGTATGCA
138481   CATGAAAAGA TTTTAAACAT CATTAGTTAC TAGGGAAATG CAAGTCAAAA CCACAATGAG
138541   ATACTTCACA TTCAACAGAA TAGCTAATGT TAAAAGGACT GACAATCCCC AGGGTGAGCA
138601   AGGGTGTGGA GGAAACTACT CTCATATATT GTGAATGTAA GAGGACAATG TTACAACTAC
138661   TTTGAAAAAA GTTTGGCTGT TTCTAACATA AAATTAAACA CTTATACAGC CCAGCAATAT
138721   TTCTGGGTCA TTTCTCCCAG ATAAATGAAC ACATGTCCAT ACTATGACAT GTACAAATGT
138781   TCATACTGGC TTTGTTTCAC AATGCTATAA ACTGGAAACA ACCCACGTGT CCATCAACAG
138841   GTGAATGGGT AAATAAATTG TAATATATCG GCCAGACGCA GTGGTTCATG CCTGTAATCC
138901   CAGAACTTTG GGAGGCCAAG ATGTACGGAT CACCTGAGAT CAGGAGTTTG AGACCAGCCC
138961   ATCCAACATG GTGAAACCCC ATCTCTACTA AAAAATTAGC TGGGCATGGT CACGGGCGCC
139021   TGTAATCCCA GCTACTCGGA AGGCTGAGGC AAGAGAATCA CTTGAACCGA GAGAGGCGGAG
139081   GTTGCAGTGA GCCAAGACCA TGCCATTGCA CTTCAGCCTG GCAACAAGA TGGAAACTCC
139141   ATCTCAAAAA AAAAAAAAAT TGCAATATAT CTATATCTTG GAATATTATA AAGCAATAAA
139201   AGGGAATAAA CTACTGATAT ATACACAAAA TGGATGAATC TCAAAAATGT GAAGGAAAAT
139261   AAAAAATACA TATGATATAA ATTCCATTCA TATGAAATTT TAGGAATGGG AAAACTAAGC
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139321   TGTAATTATG GAAAGTACAT CAGTGGCTGC CTGGGGCCAA GAGGATGGAA GAGGCGGCAC
139381   AGGTGATACT ACAAATGGAA ACTATCTAGG TTGACGGAAG TGTTCTGTAA CTTGATTACA
139441   GTAGTAACTG TTTGGGTATA TAAAACGCAT CAAATTGTAT AATTAATACA GGTGTATTTT
139501   ACTGTGTATA AATTATTCCT CAATAAAGTT GATTTTTCAT TAAATATATT ATTTGCTAAA
139561   ATGAGGAGAG ACAACTATTA TCTTAAAATA GTTAAGCACA ATAAAAATAC TACAATCAAC
139621   TCATTATATA TGGAAATTAA AGGAGAAAAA TAGTGGTATG ATTAATTAAA ATAAAAAGAA
139681   AACCTTCTAA ATTTTATCTT AGCTCATAGT TGTAAAAGCT GCCATCCCTA ACCAAGGCCA
139741   CCCTTGACCC TTTCTCATGT TCCATCTTTC TGTTTGTTTC ATAGTTTATG TCTCACCAAA
139801   ATCTATCAGA TAAACGTATT CATATGAAGA TTTAAATATA TTACATGTTA AGCCTTAGCG
139861   AATACTTCAA TATCTAAAGA AGGTACAAAC AAAACAAAAA TCAACACTTA GTTATAAGAG
139921   ATTACATACT CTCCAGGGAA GACCTGAAGA CTAGCCCCTT TCTGGATCCC ACTAGCCCCT
139981   CATCCCACTC CAAGCCCTCC CCTCCAATCC CATATGCACT GGGCATTCAT ACAAATAAGA
140041   CCATCAGCTC TGGATATCTG TACTGATTGA TGCTCCTGCT AACTACCTGA ATGATTGCGA
140101   TGTAAGGACA GCACTGCCTG AATCCTATTT ATCTCTCGCT ATGCCATAGC GGCCTTCCAT
140161   GCTGATGGCG TGTTTGAGGA TCCAGAGGGG TCTTTGGTTG GCAGGATTGT TTTATTTCCC
140221   CAAGAGGAGA GCCTTGATGC AAAAATAGGT GAAGAAATCA GTACAACAAA ACAGAAAGCC
140281   TAGAAACTAC TATGAACACA ATAGAGCAGA AGTAGCCTTA AGAGTTGGTG GAGAAAGGAT
140341   GGTCTATTCA ATTACCTGGG CTGAGAAACT GGCTTTCATA TGGAATAAAA ATAAAATTAT
140401   AGCTATACCC CATATCATAC ACAAAAGTTT CTACATCTAA CAAAGACACA GATAGAAAAT
140461   GTTTTAAAAT TTTAGAAGAA AATAGTGCAG AATTTTAGTG CAGAATTTCT TAGACTAGAT
140521   GCAAAAACAA AAATGATTAA AGTGGCCAGG CACGGTGGCT TATGCCTGTA ATCTCAGCAC
140581   TCTGGGAGGC CGAGGTAGGT GGATTAGTGG AGGTCATGAT TTCGAGACCA GCCTGGACAA
140641   CATAGTGAAA CCCCATCTCT ACTAAAATAC AAAAATTGGT AGGGTGTGGT GGCTCACGCT
140701   TTTAATCCCA GCTACTTGGG AGTCTGAGGC AGGAGAATCA CTTGAACCTG GGAGGCAGAG
140761   GTTGCAGTGA GGGGAGATGG CGCCACTGCA CTCCAGCCTG AGCAACACAG CGAGACTCTG
140821   TCTCAAAAAA ATCTAAAAAT AAAAGATTA TTTTTAAAAG ACTATTTTAA ACAAAAAAAA
140881   TCGTTTAAAT GATATGACAC ACTACATCTA ATATTTGGAA AAGTACTTCT TAATACTTTT
140941   AATAAAAAGA GGCGCTGAGA GCATACAACC TATCCTCAGA AGAGTGTTTG ACCTCTAGGA
141001   GGGACGCAAG CGCGTTCTTC CTTCATTTTA ACTGGTCATT TTCATTTATT TCAGGAACAT
141061   CTGAAGTAAA CACAGTCACA CGTTAACCTT TAAAAATCTA GGAGGTGCGT ACGCATAGTT
141121   CCATTACTTC AATTTTTGTA CTTTTGCATT TTAAAATATC ACAGGGAAGC TCGGTACAGC
141181   TTCAAGGCTA GGAGGGGTGG CTCTCTCTTA AGCCCTGTCC CCGCCAGCCC CAGACCTCTC
141241   GTCCCGCCCC CATTGCCCAG TCCCCACCCT CACTTCCCCA TTTCCCCACT CCCGCGGTCT
141301   CTTAACGCAC CTCGTTTTTC GTCCAGTGGA CTCAGACCTG TAGTCTTCCA CCAGGATCGG
141361   CTCCTTTCCC GGAGCTCTCG CTCTTAGAGG AAATTGAGAG AAGCATCAGC GGAGACCCAT
141421   CTGTGGCTCT CCAGAGGGCG CGGCATTCAG ACCCCAGATC CAGCTGTGAG AACGGACCCC
141481   AGGCTCACAC CAGGCCTGCG GGAGGCGGCC CACCAGAGGC GCTAGAAAAC AAGCCTCGCG
141541   GGGAGGCGCG CAGGGCGACT GCAAGCTGTA GGGGGCGCTG GCGCCCTCAC AGGCCAGGGG
141601   CAGGGCCGGC GCTGCGGGCG GGGCTCCTGC GGCGTGAGGG GCGGCCCCAG GCCAGCAGCT
141661   GCGCCCTGGC TGGGAGCCGG GGAGCATTTG CTGCTCTGCT GGACCCTGAG TCTGGCGGCG
141721   GGCGGCCTCC TCTCCGCTCC CCGCCCGCCA TCCCCCAACT CCCGATCTCT CTGCTGCGTC
141781   TGGCCTCAGG CTGAGACCCC AACGAATCAT TCCCCGCATG GGAACATTTT ATGATATAAC
141841   TGAATTCAGT TTTATGTATA ACTGAATTAC GGATATGAGA ATCTCAAATG AGGACGAATG
141901   GTTTTTACGC ACAAAACATG AGACACAAAT CTGTAAGAAA TATAAAGTCG TGACCACGTC
141961   CTTTCAGAAC TTTAACCTGT TTGCTGAAGT ACGTCAGTAA CAATGGCAGG GAAAGGGTAT
142021   CTTAAATTTC ACCACAGCCT CAAAGAGGCC ATTTCGTGGA TCCGCTGAGG CTTGGAGTCG
142081   GCCTTCTGAC CACGAGTCCT GCGGCTATGA AAGAGGAAGC CGCGGTTCAG GGCGTCCTCG
142141   CGAGTCGTGC AGCCCGCCCT GCTCCAGCTG GGGACACCGG TGGTCACGGC GCTTTCCAGC
142201   TGCAGATCCA GGCGGCAGCC CAAGATTTGG TCCAGCCGCC AAGGGGTGGC TCGAGTGACT
142261   GACGGGCCTT GAACGCTCCC AGGACCCACA TCTGGAGAGG GAGGTGGGGG TGGGGTGCTG
142321   AAGTCATTCT TGGGGCCCCT GGGGGCGGGC ATGGACCTGG GTAAGGCCAG AGAAATTGAC
142381   ACCTCGTGAC ATCCCTGGAA GAGAAGTACG TTCAGTGTCA CTCCAGAGCT GAAACCGCCT
142441   TCTGGCTGGT CCCTCCTCAC CTACATACTT TTCTAATTTG TCTGGAGCAG GCCGGGCATC
142501   TGTATTATCT GGTTATTTAA ATATCTGGTT ATTTAAAAGC TCTCCATTAA ATTCACATAC
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142561  ACGAAAATAA  AAAATTAAAAA  AAAATTTTAAA  AAAAAGAAAC  AAAAGCTCTC  TAATGACCAA
142621  GTCCTACACG  ATAGTGAATA  AATTTTTTTG  TGTGGTCCCT  AAAATTGAGT  TCATGCCTTT
142681  TCTGAAGTAA  TAGACGCCCA  GAGAAGGGAT  CGACTTACCC  ATCATGCCAC  AGAGATTAAT
142741  TGGCCCCAGA  ATTCTTTAGC  AGACCGTGTA  TATGAACGTC  CTTTGCAATC  ATATAAATTA
142801  ACTGGGAAAA  CCTCATTTAG  TATGTTACAT  GCCTAGCGTT  TTGTGCCTGA  ACACCTTACA
142861  AGAACCAGGG  ACTATTGCCC  CAATATTATA  TTTCAGGAAA  GGAAGGCCCA  GACAAATGGT
142921  GTCACTGGTC  CACTTTCACC  CAGTTGGTAA  ATGAAACCAG  AAATTATAGC  TGTACCACAG
142981  AAAGGTGAAA  ACGTTTCTTT  TATAATTTCA  CATACAATCT  TTAATGGACC  CAGTGTCCAA
143041  CACATTAAAG  CAAGTGCTCA  GGAGTGACAT  CAAGATGTAA  AAAATAGTCC  TGTCCTCAGG
143101  GAGTTTAGGT  CTTGGAGAAA  AGAGACCCAA  GGAGACACAA  GACAAAGGGG  AAAGAGAAGG
143161  AGCGCTGAAG  ACTGAGGACC  CTGCCTGTGG  ACTGAAGTGA  GGATGGGGAC  ACCCGATGCC
143221  CGGAATATGA  CAGTTTGGAG  GGGCCTGAAG  GACTCTTCTA  TTCTCTATCA  GAAAAACAGA
143281  ATTACTCTCC  TAACCAGAAA  AGGTATTTCA  ATTTATATTT  TCCATCACAG  CACTTTTCTG
143341  GTGATAATTT  AATGTGTTTT  AAAAAATGTA  TCACAGTGAT  GGCCTGGTGT  GAAATAAATA
143401  ATAAAATTTT  AAGAATTAAA  AAATATAAAA  ATCTTTTATA  TAGACATTAG  GAGTTACAAG
143461  GATAACTGTG  AATTATAATT  AGTAATTAAA  TTGAAATACT  GATTATTTTC  ATTTTTATTT
143521  AATTATTTAA  TAAAACCTAT  TTAACATTTA  ATATTTATCA  GTAATTAAAT  CTAATTGTTA
143581  ATATTTATTA  TTATAAATTA  TTTTAGAATT  AAAAATAAGT  GTAGAAGCGA  GGCATGGTGG
143641  CTCAAGCCTG  TAATCCCAAC  ACTTTGGGAG  GCTAAGGTGG  GAGGATTGCT  TGAGCCCAGT
143701  AGTTCAAGAC  CAGCCTGGGC  AACATGGAGA  AACCCTGTCT  CAATACAAAA  AAATGAGCCA
143761  TGTGTGGTGG  TGCGTGCCTG  TAGTCCCAGC  CATTCTGGAG  GCTGAGGTGG  GAGGATGACT
143821  TGAGCCTAGG  CAGTCAAGGC  TGCAGTGAGC  CCTGATCTTG  CCACTGCACT  CCAGTCTGGG
143881  CAACAGAGCA  AGACCCTGTG  TCAATATACA  TATGGACAAA  CTTAAAATTT  AAAATGAAAG
143941  CATACTACTG  ATACAGAATT  GAGTAGAGAT  GCAAAGCTAG  TCCTATAACC  AGAACAATAA
144001  AGATAAAAAG  GAGAGTGGAA  GAAGGTATGT  CATGAATTTC  ATGATAAATG  GCAATTGCAA
144061  ATATCCTGTA  GCAGAACAAA  ACAACAAAAC  TGTAGATAAA  ACATATCCAA  CCCTTTGGAA
144121  GGCCAAGGAG  GGAGGATTGT  TTGAGCCCAG  AAGTTGGAGA  CCAGCCTGGG  CAACATAGTG
144181  AGACCCTGTA  TCTAAAAAGG  AAGAAAGAAA  AAAAAAAAAA  GGATGATAAA  GTAGACAATA
144241  TTGAAAGCCA  TTTTCTGCAA  ATACATAGTG  AATTTGATCA  GTAATTTTCT  TCCAACAGTG
144301  CAAAAATGAA  TAGATATTAG  TTGCCTGAAA  TAAAAATCAA  ATATCCAACA  AAAAATATTG
144361  ACTATCTAAT  AGTATCTAAG  CTAGTAAATT  TGGCCAGTTA  TAAAATGTCT  TAAATTTTTA
144421  TTTAAAAAAA  GAAAACCATA  TTTATAAGAA  GAGGTGATAA  AGAGAAATTA  TTTCAGTTAT
144481  GAAGATTTTG  TTAGAAAACT  ATGAGAAAAA  AACTATTTTT  TGTTTTCAAA  AAGTGAAAGA
144541  TTAAGTTACC  AAACAGTTGC  TAAAGAATAC  CAGATGGCTG  AGCGTGGTGA  CTTATGCCTG
144601  TAATCCCAGT  ACTTTGGAAG  GCCAAGGCAG  GAGGATCATT  TTAGGCCTGG  AGTTCGAGAC
144661  CAGCCTGGGC  ACTGTAGCAA  GACCCGTCTC  TATTAAAAAA  AAAAAAAAAA  AAAAAAAGA
144721  ATACAAGACC  TTGCTAACAA  TAGCAAAGAT  CAATTAATTC  AAAATTTGAA  AAACTGTAAT
144781  TTATTTAGCT  TTAGAGTACT  CTCGTGATAT  GAGATTGCCA  AATTAATACT  TTGGGTGCAT
144841  TTCTTTTCTC  AAAGGACTTG  CAAATTTACA  AAGAAGTGTT  GAAGAAAAGC  CACACATTGG
144901  CAGGTAATGT  TTGCAAAAGA  CAGATCTGAT  GAAGAACAAT  ATTTTTAGAA  TATACAAAGA
144961  ATACTTAAAA  CTCAACAGTA  AGAAAATAAC  CTGATTTAAA  GCAGGCCAAT  GACCTGAACA
145021  TCTGTTCACC  AAAGAAGATA  CACAGATGCA  AGTATGCATA  TGAAAAGATG  CTTGACATCA
145081  TGTCATTAGG  GAACTGCAAA  TTAAAACAAG  TAGATACCAC  TGCATACCTA  GTAGAATGAC
145141  CAAAATTTAG  AACACTGTCA  GCACCAAAGG  TTGCAAAGAT  ATGTAGCAAT  AGTAACTTGT
145201  TCATTACTGG  TGAGAATGCA  AAATGTGCAA  TCACTTTGGA  AGACAGTTTG  GTGGTTTCTT
145261  ACAAAAGTAA  CCATACTTTT  ACCATAAGAT  TCACCAATCA  CACTCCTTAG  TATTTATCCA
145321  AAGGAATTGA  AAACTTATCT  CCACACAAAA  ACCTGCACAT  AGATGTTTAT  AGCAGCTTTA
145381  TTCATAATTT  ATCCAAAACT  TGGAAACAAG  ATGTCTTTCA  GTAGGTAAGT  GGATAACTGT
145441  GGTACTTCTG  AATAATGGAA  TGTTATTTAG  AGTTAAAAAG  AAATGCATTC  ACTTTGGGAG
145501  GCCGAAGTGG  GTGGATTGCT  TGAGGCCAGG  AGTTTGAGAC  CAGCCTGGTC  AACATGGGAA
145561  AACCCCAATT  AGCCGGGCAT  AGTGGCGTGA  GCCTGTAATC  CCAGCTACTC  GGGAGGCTGA
145621  GATATGAGAA  TCGTTTGAAC  CTGGGAGATG  GAGGTTGCAG  TGAGCCAGTG  CCACTGCACT
145681  TCAGCCTGGG  CAACAGAGCA  AGACTCCTCT  GTCTCAAAAA  AAAAAAAAAA  AAGAAAGAAA
145741  AGAAAAAAGA  AAAAGAAAAA  GAAAGAAAC  GATCAAGCCA  TGAAAACACA  TGAAGGAAAC
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145801   TTAAATGTAT GTTACTAAAA AGCCAACCTG AAAAGACTGC ATACTATATG ACTCCAACTG
145861   ATGCAGGGCA AGCAAGCCAA AAATTAGGGC TTAGCCCGGG AAGAATTCAA GGGTGAAGTG
145921   GTGGTGTTAG CAACTTTTAC TGAAGCAGCA GTGTACAACA GCAGAACAGG TACTGCTCCT
145981   TGCTGAGCAG GGCTAACCCA TAAGTAATGT GCCCAGAGTA GCAGCTCAGG GGCAGTTCTG
146041   CAGTAATATA CCTGCTTTTA GTTAAGTGCA TGTTAAGGGG GATTATGCAG AAATTTCTAG
146101   AAAAAGAGTG GTAACTTCGG AGTAGGTACA GAGGAAAGAA GTCGATAATG TCCTGTTGTT
146161   GCCATGGCAA CGAAAAACTG ACATGGCGCT GGTGGGCGTG TCTTATGGAG AGGTGCTTTA
146221   ACCTCGTCCC TGTTTCGGCT AGTCTTCAAT CTGGTCCGGA GTAAAGTCCC TGCCTCCGGA
146281   GTTCACTCCT GCTTCCTGCT TCACAACTGT ATGACACTCT AGAAAAGACA GTAACTATGG
146341   ACACAGTCAA AAGATTAGTT GATAGAAATT GGGTGACAGG AAGTGTTGAA AAGGCAGAAC
146401   ACAGGATTTT TAGGGCAGTG AAACTTCTGT GATACTATAA TGGTGAATAC ATGACATTAT
146461   ACATTTGTCA AAACCCATAG AAAGCACAAC ACCAAGAATA AACCCTAATG TAAATTACAG
146521   ACTTTCGTTG ATAATGACGT GTCAATGTAA GTTCAATTGT AATAAATGTA CTACTGTGGT
146581   GCTGGATGTC TATGGTGGGG GGACATTTTT GCTTCAATAG TTACAGTTGA AGTAAATGTT
146641   TGTGTTTCCC ACAATGCATA TGTAGAAACT CTCACATTCA ATGTGATGGT CTTTGGAGGT
146701   GGGCTCTTTG GGTGATAGTT AGGTTTAGTT GAGATCCTAG CAGATCGAGT CTTCATGATG
146761   GGCATGATGG GACTGGTCCC TTATAAGAAA AGACCAGAAA GCTAGCTCTC TCTTTGCCAT
146821   GTGAAGACAT AGCAGGAAGG TAGCCATCTG CAAGCTAGGA AAGGGCCTTC ACAAAGAATC
146881   AACTCAGACC TCAGAACAGT GAGAGATAAA TTGTCGTTGT TTAAGTCACT CAGGCTGTGG
146941   TATTTTGTTT CAGCAGCCCA ACCTAAGACT GTTAATTGGA TTAGAAATTT CCTTTTGGGG
147001   ATGGTGTGTG GCGGGCGGGG GGCGGGGAGT ACCTTTGTTA AGCTTTTATA TCAATGAGTT
147061   TGTAGGCTTT TCTTTTTTGG TCATTGACTA GGACAGTTTA AATAGTATGA GTGTGAAGGA
147121   GATTGTTGGT CATCTATTCG ATGTCCCTTC TCTGTTTTTT AATATGAGAA CTCCTGATTT
147181   TCAGCCAACT ACCCTGGAAA AAAAGCTAAT CTTTCTGACT TCTTAAGTGT GGCCATGTAC
147241   TAAATTCTGG CTAATGCAAG GCAAGCCAAA GGTTTTATGA TAGGTTTTAG GACACTAGAG
147301   TAAAAGAGAG CTGTTGCACA CATGCTCTTC ACCCTACTTT TGTGTCCTTT TTTCCATCCT
147361   ACAACTTGGG TTGTGAGTAT GATGGCTGGA ACTTTAGTGG CTCTCTTGGA TCCCAGGGGT
147421   AATTGAGGGG TGGCTGGAAG GAATCTGTGA TTTTCTGGAG TTTCCATACA CAAACAAGAC
147481   CTGGATTTTC TGGGCTTCCC AGACTTCCAC ATCTAGACTT GCTTTAAATG GGAGATAAAT
147541   AAACTTGTTT CAGCCACTGT CATTTTGGGC TATTTTATAG AACTTAATCT AATCTTCAAG
147601   GGTACATGAA TTGCTTTTCC TTAAAAAAAA AATCAGCCAT AAAATCATCT TCTTTTTTCT
147661   TTTGTTCCCC ACATTATTTA GTTGGAGCTC TGTAACTTTT TTTTTTTTTT TTTTGAGAC
147721   AAGGTCTTGC TCTGTCACTT AGGCTGGAAT TCAGTGGCAT GACCATGGCT CACTGCAGCC
147781   TTGCCCTCCT AGGCTCAAGC AATCCTCGTC TCAGCCTCCT GAGTAGCTGA AACTAAGGCA
147841   CATGCCACCA TGCCCAGCTA ATTTCTTTTC TTTTAGAGAT GGGAGCCTTG CCCAGGCTAG
147901   TCTCAAACTC CTAGCCTCAA GTGATCCTCC CATCTCAGCC TCCCAAAGTG ACAGGATTAC
147961   AGGTGTGAGC CACCATGCCT GGCTGCTCTG TAAGTGTCTG AATTTCATTT TGTATTTATC
148021   AGTCTGTTTA GATTTTCTTT CCCTTCTTGG GTCAGTTAGG CCATTGGTTT CTTTTTAAAG
148081   GTTTTCAAAT TTATTTGCAT CTAATTCTTC AAATTACTCT CAAAATTATT CCAGTATATA
148141   TTCTTTTGTT CCTATTTTCT TCTGTATTCT TTATTAAAAT AGCTAATGAT TTATCTAGCA
148201   GGACTTATAT TCTTTCCATA ACTTTCCTGC ACCCCAATTA ATCTCCAATT TTATATTTCT
148261   TCTGGCCTTC CTTATAGTTT CCACAGGTTT ATTTTATTCA TTTTTTAAAA CTTTTATTTA
148321   ATTGTTTATT TTATTATCAT TCTTTCTTAT TCAGCAATCT AAGTGCTTAG GGATATAGAA
148381   TTTCCTCTAA GCAGCATATG CTAGGCTTTA ACAATGTTAG GGAGGCCTCC CCTTTCTGGG
148441   GAAGACCACA CTTACATTAA CACAGGACTG TGGGATGCCA AGAGGTAGAG AAGAGCTTAT
148501   GAATATCCAG ATTACATCTT CACTGATCCT GCACAAAGGT GGGGTTCCTC GGTTACCCAC
148561   TGGGTCCTAT TACCCAAGTC TGGGTCAGCA TACCGAGACT ACGGGTATAT AGAACAAGTG
148621   CAACTGGCGA TAATCCTTCT GTTGGGGAGA AAAATCTTTT TTTTCTATTC ATCTTAGGTT
148681   CTCCATCTGT GGCCCTATCA AGTAGACTAA CAAAAGACAG ATTGACAAGA CAGAAACAAA
148741   GCATGTGCAT TGTACAAACA CAGGGGAGTA CTGAGATGAA TACTCAAAAG AGGATTTAGA
148801   ACTTGGGCTT ATATAGCATT TTAAGAAAAG AATACATTTT TTAAGTGACA AGGAAGACGA
148861   AAAGGACTTT GAGTTTCTAG TGCAGTAAAT TGTGGGAAGG CAACTTTTTC TTTCCCTTTT
148921   TTTTTTTTTT TTTTTAAAAA AAAAGACTTC TCTGGTGCTA TGTCCAGGCT GATAAGAGTC
148981   TAAAGTCTCT GGTGACTAAC TTTTGTTCTT CCCCGAGTAA GAAGACACCT TCACAATTTC
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149041  ATATCCTGCT  TTTAGGCAAA  TAGGGAGAGG  GCAGAGGTGT  TTGTTTGTTT  TTAATCTATT
149101  TTTTTTCTCA  ATTGTCTTCA  ACTCAAAATA  CTTCTTATGC  CAAAGATGGC  ATATTCTGCT
149161  ACCCTTCACT  TACTACTTAC  AACCCAGCCT  CTATCATCAT  AATTAGAACT  TCTGACCCTG
149221  GGGAACATGG  GCAATAGTTT  GAACTCTTTT  ATATCTCCCT  TAGGCAGAGA  TGGAGGCCCA
149281  GCCATGCCTC  TGACATCTAG  ACACAACTGT  TGCTTCATTT  CTCCTATTCT  CAGAGGTGAT
149341  GTTGTAGGAC  TTCAACAAAT  ATCAGTAAAC  ATTAATTTTT  TTTTTCCTTG  AGGCACAGCA
149401  TGATCTTGGC  TTACTGCAGC  TGCTGCAGGC  TCAAGCAATT  CTCCTGCCTT  GGCCTCACGA
149461  GTAGCTGGGT  TACAGGCCCC  TACCACCATG  CCCGGCTAAT  TTTTGTATTT  TTAGTAGAGA
149521  CAGGGTTTCA  CCATGTTGGC  CAGGCTGGTG  TTGAACTCCT  GACCTCAAGT  GATCCACCTG
149581  CCTCAGCCTC  ACATAGTTCT  GGGATTACAG  GCGTGAGCCA  CCATGCCTGG  CCATCAATTT
149641  TTATGTCAAC  TCTAAATTAT  AACATTTAGC  AATTTGTGA   CTTTTTATGG  TCATCATTAA
149701  TGTTGTTTAT  GTTTTAGTTG  TAGTCCTGTC  ATTACTCACT  CGGGTATGGT  AATTTGGTCT
149761  TTTTCAAAAT  GAAGTTAAGG  TCTATTTGCT  CTTCTCTGAA  TCATAATAAG  AACTGCCAAC
149821  AGCCATTTCA  GCAATAACTA  TTTACTGAGA  TTTTAAAATA  TTTCAAGGTA  ATTGGTCCTA
149881  GCAGACTGGA  AAATACCAAA  TTCTTTTCCA  GAACTGAATC  CCCCATCAAA  GTTCAATTTT
149941  ACTCATAATT  CCCTTTTCAT  TTGAAGCATC  TCATTGTAAG  CCAGTCTTAA  CCCTTCTCTC
150001  ACACTTTGCT  TGGCTGTTTC  TCAGGTAGAA  CTCAGTAAGT  CTGGTAGCCT  CCAGGACTGC
150061  CGCTTAGATT  ATTAAACAAC  ATGTCAGTGG  TTGGAAGAGT  CAATGTTATT  TTGATTTTTC
150121  TGTTTTGTTT  TGTTTTAAAT  GCAGTTGGCG  GATAATTGCA  GCTTTCTTTC  ATTCCCTACA
150181  TGAGTTCAAA  TGGCAGCAAA  CAAACTAGGA  GAACGCAGAC  CTTCTGACTT  GTGGGTACCC
150241  CTACTCATCA  CCTGAAGACC  CTTGGAAATC  AAAGCCCTGA  CCCATTAAAG  ACGGATGGAG
150301  ACAGCAACAT  ACGATCATCA  CTATTATCTT  GCTTTGCCCC  AGTCCAGGTT  AACCATCTGT
150361  GGTATTTTTA  GTTGCTAAGT  CCATATATTC  AACATAAATC  AATTATATAT  CCACTAAAAT
150421  CTCAGCACTA  GTCTAACTAC  TAAGGAAATG  ACAGCGAAGA  AAACAGACCA  AACGTCTGCC
150481  CTTATGGGAT  TTATATTATT  TTCTCTGTGC  TGGTTAAACC  AAGGAGCTTC  TGCTCTTTTC
150541  CTTAGTCACC  TGGGGGAGGC  AGAAACAAAG  GAGAATATTG  ATAAACCTGG  AAATAGGGCC
150601  GGAGAGTATC  AGAGAAGGAA  GCCTTCGGGA  AAGTAAAGAT  GTGGCAGCCA  GTATTCCCGT
150661  TATAAAAGGA  TACAACTCCG  GCCTCATAGT  CCAGAAAAAT  TCCCACAAGC  AGGGGCTGCT
150721  CATGCAGATG  AAGGGAAGTT  GGGGGAGAAG  TAAGTGCTAC  ATAGCCTTTC  TTTTTGCACA
150781  GCCTGAGGGT  CCAGAATCCA  GACTGAGGCT  CTTGCTTCAT  GCCAGTGCCC  CTCTGCACAT
150841  TTTCCATACA  AACTCCTAAA  TCCCATCCGG  TTCCTTCGCC  AACATCCACT  TCAAAGTAAC
150901  GTCTTCCTGA  GGTGAAGCCT  TCACAACCCA  AGACACAGGG  GAAGGCAGTA  AATCTCCTGG
150961  AAGATGTGTC  CTGATTCTCC  TGGGTGTATC  CACGAGTCAC  TTGTCTCCGA  TCCTCAGAGA
151021  GAATTAGTTC  GTGATGAGCT  GTATCTGGAT  CCAGAGTCAC  ACTAACTGCA  AAACAAAACA
151081  AAACAAACAA  AAATAATTTT  GTTGCTGTGA  AGAACACAGG  TTATTTTATT  TTATTTTATT
151141  TTGAGATGGA  GTGTTGCTGT  CACCCAGGCT  GGAGTGCACT  GGCACTATCT  CAACTCACTG
151201  CAACCTCCAC  CTCCTGGATT  CAGGCAATTC  TCCTGCCTCA  GCCTCCGGAG  TAACTGCGAC
151261  TACAGGTGCG  CACCACCACA  AGTGGCTAAT  TTTTTAAAT   TTTCTGTAGA  GATGGGGTTT
151321  CGCCATGTTG  GCCAGGCTGG  TCTCAAACTC  CTGACCTGAA  GTGTTCCACC  CACCTCGGCC
151381  TCCCAAAGTG  CTGGATTACA  CAGGTGTGAG  CCACCATGCC  CAGCCACAAG  TTATTTTCAA
151441  TAAAACCAGC  CTGTGTTCAA  ACCCAACTAT  TGTTTCTTAT  AAACTGGGTG  AGCTTAGGCA
151501  AATCATTTAA  CTTTCTGAGC  CTCAGTTTGT  TAACTATAAA  GTGGAAATTA  CCGTATTTGT
151561  TGCAGAGAAT  GGTGGGTAGG  ATTGAATAAG  CTTATGTTTG  CTTAATGCTT  GGTAAAATTC
151621  CTGGTACATG  GTAACCACCT  AATAAGTGGT  AGTTGTTGGG  GTGATCAGGC  CCAACACCAG
151681  GCCGTGGGGG  CTACAAAGTC  CGGCGGGGTC  AAAGGAATGA  GAAAAGACAA  GTTAAGAGTG
151741  CATAAAGTGG  GTCCAGGGTG  CCAGCACTAG  ATTGGAGGCT  GCAAAGGCCC  TAAGCTCTGG
151801  GAGCCCACAC  TATTTATTGG  TGATCAAACA  AAGAAGCAGG  TGGTGAGGAC  GTGAGGGTAA
151861  ACAGGTGAGG  GCATGAGGAC  ATGGGGGTAG  AAAGGTAGTG  GTGCATTAAG  CGTAGCTGTG
151921  ACAGTTTAGC  ATTTTCTTTG  ACACATGTAG  AATATACTCT  GCTGCTTGAG  ATAGTAGAGG
151981  ACACGTTTAT  GAGTGAAAAG  CAAGGAACCA  ACAAGTCTGT  GCACTTTCCA  GAGGCTATGA
152041  GGGGTTTTAT  GCCCTGAGCC  CTGGGTTCCA  TCCAAGCCAC  AAGGGGTTTT  ATGCCCTAGG
152101  CTTAGATTTG  TGGTGCGGCA  GGGCAGCCTT  CCACCATTTG  GCACAGAGCT  TGGTGTTCCA
152161  AAGGCCACGA  GGGGTTTTGG  ACCCTGGACC  CCGGACATCT  TCCAAGACTC  TTTTACATTA
152221  TGACAGACAA  GCCAGTCCTG  CTTCAGCTCT  TCTAACAACA  TGTAGTAATA  ATGATATCAT
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152281  CAACATCATC TTCGTCTTAA TTATTCAAGG ATGCCAAGGT ACAGAACTAA CCTGTTAATA
152341  TGGTTACCAT CCTGTCCAAA GTTCTTCTCC CATGCAGGAC TTCCAGGAAT CATGAGACAG
152401  TTGAGCAGAA AGATACCTTT TCCCTTCTCT ACTGAATAAC CACCAACATT GAGAATCAGA
152461  GAGGGAAAAT GACTCAGCTA ATGTCTTAGC TTGTTATTGG AAGACCCAGG TCTCATGACA
152521  CATGCCTAGT CCCATGACTT TTAATTGTAA GCTCTTCTCT TTCCCCTCAG ATAATGTTCC
152581  ATAAGCATTA GTATGAGATA ATAATACACT GAGGACCAAT ATACATGAAA AATATCAGAC
152641  TAGAATCAAA CAAGACAGAA AAAAGATCTG ATAACCTAAA GTGAGATACT GAACAGTATG
152701  CAGTTTTAAA AATAAAAAAT GGTAATAGGA TGTTCTAACA AGAGAGTTAA GAAACCACTG
152761  TGCTACTGAG TTAAATGTTG ATCAGTTGGT CTGTGACAAT TAAGGAATTC AAGTATTCAG
152821  AAACACTTCC TGTGCTGGAT GCTCTCTGTT TGTTCTTCCA AATAATCCCT CACTTTTCCC
152881  TGTCTTGCTC TGTGCCCAGG AAGGCTGACA TGGACAGATT AACCAGGCTT TCCGCCCTCT
152941  GGCTTGGTTC AGCCAATGGG AAGCACCAGA GGAGACCATA GGGCACAAAG AAGCAGCCTT
153001  GGGAGTATTC AGTACCCCAG TCCCACGCTA TGATTGGAG  GGTCTGCATT CCTCTGCCTC
153061  TGGGCACACT CTAGTATAGT TACAGCTCCC TACACCTGCC ACTTGAGGCC CAGAGGAGGT
153121  GATGGCTCTC TAACTGTTCC TAGTTCTGGG TGCTTCCTGT TCCTTGTGGA TTTCCCAACT
153181  CCTCACCTTT GTAAATACCC TCCTTTTTCA AACTCTATTC AGTTAGCTTT TATCAGCCTG
153241  ACTCACAGAA GTTTGGGGTT TCAATTCATA TTACCTGAAT GACCCAGGAA AACCCATGTT
153301  GAGAAATTAA AATGTTTACG GGGTGGTAAT ACCACTTAAG AGAAAAAATA TCAATTGGAT
153361  TTTTAAAATT CCACCTATCT ATTGGTGTGA CACATCAACA AAAACATATA GAAAGATTGG
153421  AAGCTAAAAG ATAGATAATA TAGTCATATA CTGTTATAGT ATTATATCAA AAGATATTAA
153481  GTCAGAGCAT TATTAAGAAT GGAAGAAGGG CCAGGTGTGG TGGCTCATGC CTGTAATCCC
153541  AGCACTTTGG GAGGCCAAGG CAGGCGGATC ACTTGAAGCC AGGAGTTCAA GACCAGCCTG
153601  CCCAACATGG CAAAACCCTG GCTCTACCAA AAATACAACA ATTAGCTGGG CATTGTGGCA
153661  CATGCCTGTA ATCCCAGCTA CTTGGGAGGC TGAAGCACAA GAATCACTTG AACCGGGGAG
153721  GCAGAGGTTG CAGTGAGCTG AGATTCGCC  ACTACACTAC AGCCTGGGTG ACAGAGAGAG
153781  ATTCTGTCTC AAAAAAAAAA AAAAGAAAG  AATGAAAGGA GTCACCTAAA AAAGATAACA
153841  CAATTTTAAA CATAAATGTA CTACATTATT AGTGAATTCA TGTTTAGAAT TGTGTTAATA
153901  TACAAAGCAA AAATTGTAGA ATTATAGGAG AAATGGACAA ATCTACAATC ATCATGGGAT
153961  GTTTTAACAT TCTTCTTTCC ATAATTGATA GATCAGGCAG ACCAAAAGAA AGAAATAAGG
154021  GAAGATACGG AAGGTCTGAA CAATCTAAGA AGCGCAATCT CATAGTCAAT ACATAAAGCT
154081  CAGCAATTGT TTAATAATAG TAAGCAGAGA ATATGCAGTT TTCTCAGGTA TAGATGGAAC
154141  ATGCACTAAC TGAGTAAATA CTAGGCAGAA AACAGTCTGA ACAAGTTTCA ATAAATCTGT
154201  ATTACACAGA TCATTTTCTC TAGCCTCAAT ATAAGATTAT AAACCAATAA TAAAAAGATG
154261  ACTAAAAAGA TTCTAAATAT TAGGAAATGT AAACTACTAA TAAGTCATTA GAAGATGTAT
154321  AGAATGGAAC AATAATAAAA AGTTATTTAT AAAAATATAC AATGAAGCTA AAGCAGAATT
154381  TTAAGGAAAA TTTGTAGGCT TTAAATGCTT ATCTTAGAAA AATTAAAAAG CTGAACATTA
154441  ATGAGCCAAG CATCTAATTT AAATTTTAAA AAGAACATAG AAAGCCAAAT ATAATTTTTT
154501  AAAAAGAAAA AATAGATATT AAACAATATA ACAGTGAAGT TAAAGAAAAC AAGAATGCAA
154561  TAAAGAGGAA AAACAAACAA AAAAAAAGGT AGCTTCTTTT AAAAGAAATT TAATAAAAATA
154621  GACATACCTC CAATGAGATT TATCAAAGTA AGACAGAAGG CACAAATGGA ATGAATACAG
154681  AAACTTTTTA AATATTACAG AACTTTATAA TAAATCTTAT GCTACTAATA AAATTGAAAG
154741  TACTGATAAA ATTATTACTT CCTAGAAAAA ATATTTCTGA GTAAAACTCA CTCAAAAAAC
154801  AAATAAAGCA TGGGCAGACC TAACATTAAA GAAATGAAAT CACTACTTTA AATTTTACCG
154861  ACAGATAATA AAACGTGCAT CTTTATCAAG CAAAAATGGA ACTTGTCAGT TTTATAGGAA
154921  ATTTAGAAGT CAAGGCATGA GTAATGCCAA TCTCATACCA AATCCTACAA AGAATAGAAA
154981  ATTATGGCTC CCGCTTATAG ACATAGATAT AGAACTCCTG CACAAAATAA TATAAATAAC
155041  AAACCAAATT TTATATTTGC AACTATACAT ATTATATGTG TATGTATTAT ATATGTTAAC
155101  ATATACATAT ATAATATGTA TAGCATATGT TCTACATATT ATATATGTAT AGTGTATGTA
155161  TTTTACAATA TATAAATGAA AACCCAATCT TTAATATATT CATCTAGATT GTCATATATG
155221  ACATATATAA TACATTACAT CAAAAATGTG TACAATAATC AGGCCAGGCA CAGTGACTCA
155281  TGCCTGTAAT CCCAGCACGT GGGAGGCTG  AGGCGGGTCA ATCACTTGAG TCCAAGAGTT
155341  TGAGACCAGC CTGGTCAATA TGGCCAAATT CCATCTCTAC AAAAAATATG AAAAATTATC
155401  CAGGCATTGT GGTGCACACC AATAGTCCCA GCTACTCGGG AAGCTGAGGT GAGAGGATCA
155461  CTTGAGCCTG GGAGGTGGAG ATTGCAGTGA GTCGAGATTG CGCCAGTGCA CTCCAGCCTG
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155521   GGTGGCAAAG GGAGACCCTG TCTCAAAAAA AAATTAAAAA ATTAGCCAGG TATGGTGGCC
155581   TGTTCCTGTA GTCCCAGCAA CTGGGGAGGC TGAGGTGAGA AGATCACTTT AGCTCAGGTG
155641   GTGGAGCCAT GATCGCACCA CTGTACCACT CGGCTTGGGC AACAGAGTGA GAGCCTGTCT
155701   CGAAAAAACA AATATATACA CACAGTAATC AATATATATA TTATATGTAC CAATCAATGC
155761   TTCACTTTTA TATATAATAT AGATTACATC TTATTAGATA TATAGTATTC CTTCTCCATA
155821   GATAGATAGA TACAGATATA GACATAGTAT CCTCTATCCA TATTAGAGAG AGGATACTAT
155881   ATATATCTAT AGCATATAGA GATGCTGTCT CAAAAAAATT TAAACATCAG CCAGATGTGG
155941   TGGCCCATGC CTGTAGTCCC AGCTACTGGG GAGGCTGAAA TGAGAGGATT GCCATTGATC
156001   CTCTCATTGG TTGAGCCATA ATCGCACTAC TGCACCACTC AGCCTGGGAG ACAGAGGGAG
156061   ACCTGAGGTG GAAGGATATA GATATAGATA TATAAATAAA TATGTATAGA GAGAATATAA
156121   TATATGTGTG TATGTGTATA TATATATATT ATGAAGACAC TGGGAGAGAA TACTATATAT
156181   ATATGTGTGT GTGTATATAT ATATTATGAA GACACTGGTG GGATGGTTTC ATTACCAATT
156241   GGACCAAGAG TCCAGGTATG GAGCCAACAT GCAATGTTGT TGTTGACTGA GCTGGCAGAG
156301   CACTGGTCAT AGTTACGGGA AAAGAAGGTC TCCAATGAGA CATACTTAAC AAAATATATG
156361   AACTTGCCAT ATACGTGGAG AGTTCTGGTG TGTATATAGC CTTCTCTCAC CAACCTAGCA
156421   ATTGTCTTCA TCATCATTAT AATGCTATCA GAGCAAAGAT GACAGCTAAA TTTTTTTGTC
156481   CCTTTCTTCT TCTTTCTCTT CCTTCCCCTC CCCCACCTCT TTCTCTTCCT CCTCCTCCTT
156541   CATCTCTCTT CTTTTTTTTT TTGAGATGGA GTCTTACTCT GTCGCTCAAG CTGGAGTGCA
156601   GTGGCACAAT CTCAGCTCAC TGCAACCTCT GCCTTCTGGG TTCAAGCAAT TCTGCCTAAG
156661   CCTCCAGAGT AGCTAGGACT GCAAGTGCAC ACCACCACAC CTGGCTAATT TTTGTATTTT
156721   TAGTAGAGAT AGGGTTTCAC AATGCTGGCC AGGCTGGTCT CAAACTCCTG CCCTCAAGTG
156781   ATCCTCCTGC CTCGGCCTCC CAATGTGCTG GGATTACAGG CGTAAGCCAC TGTACCCGGC
156841   CTCCTCCTTT AATAGACAGG GTCTAGCTCT GTTGCCCAGG CTGGGTACAG TGGCGTGATC
156901   ATAGCTTACT GCAGCCTCGA ACTCCTGGGC TCAGGAGATC CTCCTGCCCT AGTCTCCCCA
156961   GTAGCTGGAA CTACAGGCAT AGCACACGGG GCTAATAAAA TTAATTAGGT GATAAAATTC
157021   ACTGCCCACT GATGACTAAG CTCTTTGGAC ATAAAAGACA CAGACCTTGA AGGAAAATGT
157081   GTCTACTTAA TTTTGAAACC CTATTTATCA AAAAACAGGA TGAAAATGCA AAATGCCATC
157141   CACATGCCAG AAGATATCAG CTATAATAAG TTCCCATAAA TCAATAAGGA AAAGAACCCA
157201   ATAAAAATTA TTAAACCACA GTAAATCATG GGTAAATCAC AGAGGCCTGA AGGGCTAATG
157261   GACATACAAA AAGAATCTCA ATCTCACTAG TGAAATCAGA AAAGCACAAA TTAAGTACAC
157321   AATTAGGTAC CATTTTAAAT CTGTAAGACT GTCAAAATCA TAAATTATAT AAGTAAAGAC
157381   TCAGGGAGTT TTGGAGGAGT GAGAGCTCTT ATATTGCTTG TGGGGTAGAA TTGGAACAAT
157441   TTCAAGATCT GTAGTATCTG GTAAAATTAT GATATGCATC CCTCACACCA GCATGTCACT
157501   CCAAGGTATC TCCCTGGAGG GAACATTTAC GGGACACAAG GAAGCATGGA TAAGAATGTT
157561   CACAGTAGTA TTGTCTGCAA CAGCAACAAC AACAAAAAAA CCCAACTACA CACAACTTCA
157621   ATGCCCAGTC CACAAGGCAA TGGATTAAAT AAACTTCAGG CCGGAGATGG TGGTTCATGC
157681   CTGTAATCCC AACACTTTAG AAGGCCGAGG CGAGAGGACT GCTTGAGCCC AGGAGTTCAA
157741   GACCAGCCTG AACAAAATAA AGAGATAGTG TTTCTACAAA AAATTTTTAA AAAATTAGCC
157801   AGACGTGGCA GTGCTTGCCT GTGGTCCCAG CTACTGGGGA AGCTGACGTG GGAGGATTGC
157861   TTAAGCCCAG GAATTTAAGG CTGCAGGGAG CCATGATGGG GCCATTGCAC TCCAGCCTGG
157921   GTGACAGAGT GAGACCCTGT CTAAAGAGA TAAGTAAATA ACAACTTTGC ATTTTCTGCC
157981   ACATTGCAAA ATGGTGAGAG AGTGGTTTCT AGACTCTAGA CTCTTTCTAT GACTACCTTC
158041   TAGTTATGAG ATCCTACAAC ACTCACCTAA CCTCTCTGTG TCATATTTCC TCCTCTATAA
158101   AGCAAAAATG CCCCATATAG AGAGGACTGT GATATAAAAC AAGAACCAAG AAAAGTAAAG
158161   CTTTTCTAAT CTGTCACAGA CTAAAGAGTG CTCAGTATAT GTGAGTCATT ATTCCTGGTG
158221   CTGGTAGGAG TGTATGTTAC AACTTTGAGT CAAGTAATAT GGTACCATAT ATTAAGATTA
158281   ACAACAACCT CGGCAATCCC AGTTTGGGGT ATGTTCCCAA AAGAAATGAA AGCACCAGGA
158341   TATAAGGATG CATGGACTAG AAAGTTATTG TAGCAACATT GTAATAACTA AGTTCTAAAA
158401   ACAGCCTGAA GCTCCATCAG TAGGGATATG GTTACATATA TTTATTATAT TCTTATGGAA
158461   TATTAGACAT AAAAAGTAAC GAGTAACATA GAAGAGACAG TGTATATATG TTACGTTTGT
158521   ACAAACTTAG GGAAAGATAT AGATCACCCT ACCTAGAGAA GTCAGATTGG AGACGGGTGG
158581   GAAAAACCTT GAACTTTCTC CTTATATCCT TTATATTGTT TGACTGATTA AAATGTATTT
158641   GTTGCATCTG CTTGAAGGCA ATGTAAAATA AAATAAACAT ACATTTAAAA ATAAAAATAA
158701   AATTTATTCC TATCACTTTT GTAATAAAGC TGGGCACAGT GACTAACACT TGTAATCCTA
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158761   GCACTTTGGG AGGCAGAGAC AGGCAGATCA CCTGAGGTCA GGGGTTTGAG ACCAGCCTGG
158821   CCAACATTGT GAAACCCCAT CTCTACTAAA AATACAAAAA TCAGCCAGGC ATAGTGGTGC
158881   GTACCTGTAA TCCCACGCTA CCCGGGAGGC TGAGGCGCTG GAACCCAGGA GGCAGAGGCT
158941   GCAGTGAGCT GAGATTGCGG CACTGCAAGC CAGCCTGGGT AACAGCGAGA CTCCATCTCA
159001   AAAAAAAATT TGAAAAAAGA AAAATTTTAA TAAACAGTGT TTAAGAGGGG AGAAATATTT
159061   AGTTAAAAGA TAAGCCCATT TAAGAAATAG TTTCACTTGA CCCGGAAGGC GGAGCTTGCA
159121   GTGAGCCGAG ATCGCACCAC TGCACTCCAG CCTGGGCGAC AGAGCGAGAC TCTGTCTCAA
159181   AAAAAAAAAA AAAGAAAGAA AGAAAGAAAG AAATAGTTTC ACTTGAACCA TATTATGATT
159241   CCTTCTGTAA AAGATGAGAG TAGGCAAATT GACTCAGTGA AATCCCAGCA AAACTTACAC
159301   AAAGTCTTGT TCTTCCTTCC TGTCATCTGT ATAGGATGAA ATACAGAGTG CTTTTGGGTT
159361   TTGTTGTTGT TTGTTGTTGT GTATTTGAGG GGAACACAGG TCTATAATTC CTTTTCTGAA
159421   ATCCCTGGAA CAAAATGGGC TTTGCCATTC AAATTAGTTT AGAAGTTATA AAGGCAAAAA
159481   AATGCATATA CTCTAAAGTT CAACCCCATC ATGGCCTAAG GCAGAGCCCT GTAATCAAAT
159541   TCATCAATAT ATCTGCAGCA AAACATTTAT TCAAATTAAG TGGGATAAAT AAAGACTTTT
159601   AAATAGTCTC ATCTCAGTGC CGTTCAGGGT TGGCCACTGT GGAAGACAGA CTCAAGGGTG
159661   GCCTTCTATG ATTCCTGCCT CTTGGTGTTC ACACCCTCGT AAAATTCCTT GTCTTTGAGT
159721   GTGAGCAGGG CTTATGAATT GCTTCTGACC AATAGGATAT GGCAAAGATG ATGGGATATA
159781   ATTTCTATGA TTACGTTTCA TTATGTAAGA CTCCATCTTG CTGGCAGATT TTCTCTAAAG
159841   AGTCTGTCTC CTGAGCTCTC TCTGAAGAAA TAACTGGCCA TGTTAGAAGC CCATGTGCAA
159901   AGAGCTGAGG GGTGGCCTGT AGAAGCTGTG GGCAACCTCC AGCCAACAGC CAGAAATAAC
159961   CAGGGCCAAA GTCCTGCAAC CATCAGGAAA GAAATTCTGC CTGCTACCTC AGTGAGCTTG
160021   GAAGTGGATT CTTCCTTAGC CTAGCCTCCA GATAAGAACA CAGCCTGACC AACACCTTAA
160081   CTGCAGCCTT ATCAGACCCT AAGCAGCAGG CCCAACTAAG CTGTGCCCAG ATTCCTGAAC
160141   CACAAAAATT GAGATAACAT ATCAGTGTTG TATTAAGGTT CTAAATTATG GTAATTTGTT
160201   TGTACTAATA GATAACTAAT ATAACCACCA AATCATTTCA GGTTAGGCCA GATTTTTGTA
160261   GCCAAATGAA TCATGATAAA ACTTTCCATT TTCAGGGGTT TTTTTGATTT TGTACTTACG
160321   GATACAAATT TGTGAAAGTA TAGTCAGCAC TGATTTAAAA AATCAAGGGA GCAGGAAACT
160381   CAGTAAATGG TTCTAACATT TTGGAATCTG TAAATTGGTT GTAACATTTG TCATCTGTGT
160441   TATCTAAGTC AAGTTCCTAA AATATGTGAA TGATAGGTTA TCATACTCAC CTACTTTTCT
160501   TGCATTGCTC TAAGAGTTGG CTGAGCTATT GATAATAAAC ACTATGATCA GATCTAATAC
160561   CATGATGTGC TATTATGATC ATGTGTCAGT CACAGGGCTA AGCACTTTGT ACATGTTGAT
160621   GCATTTAATT TTGATGATAA CTCAATGAAG TAGGAGCTGT TAATATTTTC ATTTTTCAGA
160681   GGGGGAAACC AAGTCACTTG GAGTAACATG GCTAATAAGT GAAAGAATAA GAATTTGAAA
160741   GGTTTGCACA GATAACCAGA ATGCAATGCT CATCACATTC ACTGAGCAGT GAATCATACT
160801   AACTAGAGAA AGTATGAAAG CTCTACTGAA ATTAACTAAA CAACCTCTCT GGCTGTGAGC
160861   CTGCCAAGGG ACAGGTGGTA AACTTGGTTA CTGCATAAGG CCCCTTCTAT CCACAGTATT
160921   CAGGAATTCT TTAGTGAACA TACCTTGATG ACTCCTTAAC ATTTTCTTCA CATCGAAGTA
160981   AAGCTTGGAA ACATTGCACA TAGTATGAAG TTCCAAGGAG ACAGCCTCTG ATGTTTCCAG
161041   CTTCACAGCC CAACTCCTAG AATAAGCAGA GGCGAGAGAT TTCTTCAGAG GTGCATTCCA
161101   TTCATTTCTA TATACGCACA CCCCTCCCCT CCTGCATTCA AACAGGACTT ACCTGCTCAA
161161   AGTGTCATTC ACATTCTATA AAGAAACAAA AAGAAAGGT GAGCATGGGA ACATCGGTAT
161221   TTCATGGGGC TTGTCATGCA GGGCTATTCT TCTTTGCTTT ACCCGAAGAA GTAAAGAGAG
161281   TTACCCTAGT CTTAGTCTTA GATATTGATG GATACTCAAA CAAAGTAATT CCCACCAGTC
161341   TTAGGTATTG ATGGATACCC AGATGGAATA ATTCCTACCA GCTTCTGGGA GATTCAGCAT
161401   GGCAGGATGT TTATCAACAT TTGCATCTAT TCTCATCCTT GCTGAAGTCT GAGGGCCAGG
161461   AGCTTTGTCC ATGCTCCCTC TGTAAGGACT AGCTTTTGGT GATCGGATTT CCTTCACAGT
161521   GAGCCCAGAT TAGAGAACAC TTATCATAAA GGTCCTTAGT GGTGAATCTG TGCACAGCCC
161581   TGAGACTGGG CCACTGCCAC TAAGATGGTG GTAGCAGGTA TCACACAGTG GTAAAGCAAT
161641   CATGCTATAC ACTCAGCCTT ACAGTATAGT CACCAATCCT GTTAGTTAGA ACCAGAATTA
161701   ATGGCTCCAG ATGTTATCT TCCTACAGAT AAAGCTGTAG ATTGTACCAT AACAGCTCTG
161761   GAGCAAGGGT TCTACAAGCA AATCAGGGAA AAGGTTATCA CTCATTTTGG CTGCCCCACT
161821   TCATCACCCA TCAGTCACCT AGTGGAGTAT TTCAGGAGAG AGTCAACAAC CAGGGTTCTC
161881   TGCACATGGG CCAAGGAGGC AAACAGTGGT AAATGTTATC CCGTGGTTTC ATTTGGCCAA
161941   GCTGTGTTCC CTCAGAAGTT TATTTTTCTA ATTGACATAA AGGTACCCTA TAAATTAGTG
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162001   AAGGCCAGCC  TGATGGCACT  GATGTACATC  TAAAAGAAAC  ATTACTTTAT  CTTCCCATGC
162061   TTCCTTACCA  TTCTCCTTTA  ATAGCACTAT  AACATACCTT  TTTTCCCTAC  TCCAAGTACA
162121   CAGCCTCACC  TGCAGCAATT  TCTGGGCTGA  GCCCTGACAT  TTTTCCTCCA  GTTCCAGGAT
162181   GTGGCTCTTG  AGTTCATTGC  TCTTCAGCCC  CAGACCAGCC  TCATAGTCCC  TCAGTCTACT
162241   CAGAGTCTGT  TGTTCTTCTT  TCTCCAGCCT  CCAGAGATAA  GACTTCTCTT  CCTCATGTAG
162301   GAAACACTGG  AGATTCTTAA  AGTCAGACCG  GATTTTTGT   CTCTGAATCT  GTACCTTCTC
162361   CTGGAGTCAA  GAAAGTATGG  TCAAAAGGTG  GAAGTAAACC  AAATGTCCAT  CTATGGATGA
162421   ATGGATAAAC  AAGAATGAAA  GTCTGACACA  CGCTACTACA  TGACAAGCCT  TGAAGACATT
162481   CAAGCAAAAT  AAGCCAGAAA  CAAAAGGGCA  AATATTGTAA  GACTTTGCTT  ATACAAGGCA
162541   TCTGGAGTAG  TTAAGTTCAT  AGAGACAGAA  AGTAAAATAG  TGGTTACAAG  GTGTTGGCAA
162601   GACCAGAAAA  TGGACAGTTA  TTGTTTAATG  GGTAGTGAGT  TTCAGTTTAG  AAGATGAAAG
162661   ATGAAACTGA  GTTGCAGTTT  GGAGATGGGA  ATGGTGATGG  TTGCACAACA  ATGTAACAAT
162721   GTAAAGCAC   TTAATTCTAC  TGAACTATAT  ACTTAAAAGT  GGTTAAATGC  TTAAGTGTTA
162781   TATATATTTT  CACACAAACA  CACACACACA  CACAATCAGC  CACTGGGACA  TTATTTTCTC
162841   ATGAGTCACT  GAAGCTGGAA  GAATGTCCCC  AGTTCCTGC   TGCAGAGTCA  TGTGTGGGAG
162901   GCAGGCACTC  AGATGTGGAA  GAGGTTGCCT  CAGATTCCTT  ATAGTCACCC  AATTAATTTT
162961   CTTGTTCTTC  AGCCAAGACA  CAGGAGAAAG  CTGGGTTAGG  AGTGCTAGAT  AATTTAATTG
163021   TGAAACTAGG  GCCAAGTTCA  AACACTTTAT  CAGTTACAAG  GATAAAAAGA  GGTTTTACT
163081   TATGATTTAA  GAAGTTAGAT  TTCTGAGTTG  GAGCGATTTT  CTTGAAGTAA  AAGCTTATAA
163141   TGAACATCAC  CCAGACTGGA  TTTTAAGACA  ACCAGGCTGG  TAAGAGGGTC  CATAATTCTT
163201   GGCAGGGGGA  GCTTTGAGTG  TGACAGGCAT  TTATTATGGT  TAACTGAGAA  ATACTGTTCT
163261   ACTACCCTAG  GGTCATCTTA  AGCATTCCTA  TGTGTAAGAC  TGACAGAAAT  CAAGTGAAAC
163321   TCTCATCTGA  GGAGATGTAA  AGTTGCAATT  TCCATTAGTG  CTGTCTAAAT  TAATGCAGTG
163381   GGAGTGTGTA  TTCAGGGCAA  TTTGAATCTA  TGTTCTTGGA  TTGCAGTCTT  CAAACTTGGC
163441   CCAAATAAAC  TCTCTACTTA  TCTTAAAAAA  ATAAAAATTA  AAAAATAAAA  ATAAATTCAT
163501   ACAGTGTTTT  GATGACTATG  ATATAGAAGA  AGGGTCTTTG  ACTTAGGATG  AGGTGGAATT
163561   TTTGTGTAGG  AGACAGGTGC  AGCTTTAACT  CTTGTATAGA  CGGGTTTTCA  TATATGTTAG
163621   TTACAATCAA  GGTCTTCCCC  ATTGCCCAAG  ATCCTAGAAA  TGGGGGAAGT  AAGAGTGTAC
163681   TCAGGAGCTC  AAGAGCAACA  TCCACAAACA  AAGATCAGGG  TAGAGGTTAG  AGAGGACTCC
163741   TGAAAGAGAG  AAAATTGGTA  ATCAGCTTGT  GGGATTTTAC  TGCAAGCTAG  TGAATTATAT
163801   AAATATAAAG  ATTGGTGCAA  AAGTAATTGT  GGTTTTTGCC  TTTACTTTAA  TGGCAAAGAC
163861   CGCAATTACT  TTTGCACAAA  CCTAAATATT  TCCATAAAAG  AATGTGGCTC  TGATAATGTG
163921   GAGGTTAGTC  AGCCACGGAA  ATAATCTGAA  AGTTTGTAGT  TGCAAGTGTG  TAGGTTGTTG
163981   CATTACTTGT  GATGTACTTA  TAAATCAAGT  ATAGGCCGGG  TGCAGTGGCT  CACGCCTGTA
164041   ATCCCAGCAC  TTTGGGAGGC  TGAGGTGGGT  GAATCACGAG  GTCAGGAGAT  CAAGACCATC
164101   CTGGCCAACA  TGGTGAAACC  CCGTCTCTAC  TAAAATACAA  AAAATTAGCC  AGGCATGGTA
164161   GCACATGCCT  GTAATCCCAG  CTACTCAAGA  GGCTGAGGCA  GGGGAATTGC  TTGAACCCGG
164221   GAGGTGGACA  TTGCAGTGAG  CTGAGATCGC  ACCACTACAC  TCCAGCAAGA  CTCCATCTCA
164281   AAAAATAGTA  ATAATTTAAA  AATAAATAAA  TAAATAAAGT  ATATTCTTT   CATCAGCTTC
164341   ATGAGCTAGA  GTAGTATGAA  TTTCAATCTG  GAGTGATCCT  GTTTTCTAAG  TGTTCACAAA
164401   GCTTGGTTTC  TGTACCTGTA  AAGTTGAGAG  CCAGATGCTC  CACTGTGGTA  AAAGTGCCAG
164461   GGTAATGAGT  TGAGGCCTGC  AAACCAGGTT  TATTTTGACG  TATTTAAAGT  TTGAGACCCA
164521   CTCGATGCTT  TTTCTAGGTA  AATAGTCATA  CTAATTCTGC  TTCTTCTGAC  TGAAGTATCA
164581   GGAATCCCAG  CCAACTACAG  TTTAAAGATG  GAAAGATTGG  TGCTAAATAC  TCATGGATGT
164641   AAACCTGGAA  CCAGGGGCAT  AAGTACAAAT  AATGGTTTCT  TCCTTGGGTT  TCATTTTTTC
164701   AATCTGGTTT  AGTGAGAATA  AATCCTCATT  GTGCTTTTCC  TCAATCATCC  CCTATGCCTA
164761   AGCTCTAGAA  TGGAAAATAG  CTTGAGATCA  ATGAAGTCAG  ATTCTTACTT  TCCATTTAGT
164821   TATTCGCATT  GCTGTGGACA  GCTTCTGCTC  CGTACATCTG  TCTTCAAGTT  GCTTCAGTTT
164881   TGTCACAGCT  TTCTGGAGCT  TTTCCTGAAG  GAAAAATTTG  ATAAGTGAAG  CCTATTCAAT
164941   TTGACTCTTC  ATTAGGGACC  TAGGGGGAAT  CCCAATCTTC  TAAGATATAT  TTGAATAATA
165001   GTGAATATTT  ATAGAGTCCT  CATTGTTTTT  TGCTAGAGAG  CATGCTAAAG  GCTATATGTG
165061   CAGGAACATA  CTGATCCCCT  TGGCAACCCT  GAATAGTTGG  TAGGATTTTA  AACTTCATTT
165121   CTGTGCTGTA  GAAAATGAGA  CTAAGAAAGG  GGTAAAATAA  CTTGCCCAAA  GGGCTATGAC
165181   TGCCAGGTGG  TGGAGCAACA  ATTGCAATCT  CATCTGCTGA  CCCAGAGCCT  GAGCTATGTC
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165241    CACCACTAGA GTCCTGCCAG GAAAAAGTTG GATATAGAAC AAGGTAATCA TCATCTAAAA
165301    GATTTTGTAA AACAACATGC TGAACCAAGC AAAACCAATA CCAGTGTTTG GCACACATGA
165361    AATTTTGTGT CTTATGAGTC AGGAAAAATC AGGATGCCAG CTGGTTATTA GAAACAGTTC
165421    ATGGAAGAGG GGAATTCTGG TATCTTTTGA ACAATGGTAT CATGAATCCA ATTTAAAATG
165481    ATTTAGTATT CATGTCAAGC TTTTAGCTTA TTCTTCAAAA CAGTTTCTCA TATTTCTATT
165541    GAAAGTGATT TGAAGCTGAC CCAAATTGCT AATTGTAGTC AATGCTGAAA GAATTGTCTC
165601    CTGTCCTCTG TAAACCCAAC AAGTATACTC ATTCATTCTC GAGTGTTCTC AGGAAAAGGT
165661    TCTATGTAAC TGTTTTAGCA AAAGATGACA TTGTCCTTAC TATATGCCAA GTGCTATTCT
165721    ATGCATTCTA TATTTTAATG TCCTCAAAGC TTATAACCAC CTCCTGTGTA TGTGTTTTAG
165781    GGAGGGAGGA CACTGCTATT ATCCCCATTT ACAGATGGAG AAACCAAGGT GTGAAGACAT
165841    TAAGTAACGT GCCCAAAATT GCCCATCTAG TAAGTGACAA AACTCAATTT CAACATAAGC
165901    TGGTTCCTTT TCTTACTACT TGGTGGAAAA GTAATTCAAA TGGGAATATG ATCATCGCAG
165961    TTATTAGCTG CTCCATGGAG TTTAAGGAAG AGCTGCCATG AGCTGAGTGG TGGTCATGAT
166021    TGACATGTCC TTAGAAGGAC TTAGAGCCTT CATACAAGAC CACCTCTGCC TCATGGAGGA
166081    CAGAATAAGG AGCCTGACAC TGGAGACAAC ATTTTCCTCA AATTTAGGCA GGACAGAGAA
166141    GGAAAAAGGA CATCAGGACT ATGCCCATTC CTCCATGCTG CCAACAGCAA AGTCCCACCT
166201    TCCTTAATAT GCTTTCTGGC AAGAAATCTG GATGGTACAC AAAACCTCTC CCTCTGCTTC
166261    ACCTTCCACA ACCAAGCATT TCCAAATCTT TGACTCTTCT TCCTGAATCG TGCTTAAAAT
166321    CTGCCCTCTC CTCCCTTTCT TATACGGATA GTTTGAATTT TACTCCTTGA TATTCCTTTT
166381    ATCATAGACA TGCCACAGTA GCTGGGCACA GTGGTTCATG CCTCTAATCC CAGCATTTTG
166441    GGAGGCTGAG ATGGGAGGGA GACCAGGGGT TTGAGGCCAG TATAAGCAAG AAAGGCAGAC
166501    CATGTCTCTA CAAAAAATAA AAAAATTATC CAGGTATGGT GGGGCATCCC TGTAGTCCTA
166561    GCTACTTGGG AGGCTGAGGT GGGAGGATTG CTTGAGCCCC AGAAGGTTGA GGCTGCAGTG
166621    AGCCGAGATT GCACCATTGT ACTCCAACCT GGGATACAGA GCAAGACCCT ACCTCAGGAA
166681    AAAAAAAAAA AAAAAAAAAA AAAAGTAGAG GTACCAGAGT GATATTTTCA ATGTCACTGA
166741    CCCTTCATTC CCCAAATGAA AATCCCCCAA TAGGTGTTCA ATTTTTACGT GTCCTTCAGG
166801    AGTTACTTCT AAGATGAACC ACTCTCTACC CTAAATGTCC CTCCCCACCA CCAAAACCAG
166861    GGACCTCCAG GCAGACATTT TTGATGGTTT GTTTTCTTTA CTAGACTGTA GATACCTAAA
166921    AGGTGATGGG TCTTTCTTCC CTGTTTTCAG GCCCTACTGC ATGGCTTTAC ATATTGTGGT
166981    TTTTCAAATG ATATTCATGG TGTGAAACAA GAAAAAATGC GGGTGTTTGG TTTGAGAACA
167041    ACCTGTTCTA AAGCAAAAAG AAATTCATCA TAACACAAAT GGATAGAGAT AAGAGTCCAA
167101    CCATCCCATT GAAGGTCAGG ATGGACAGTC TAGATAATTG AGCAAGAAAT CATCATAAAC
167161    TATTTTTCAG AAGAATGACA TGATGAAAGC TGTATTTCCA AGTCATAATG TTAGGTTTCA
167221    AGTTAAATCA TCTCAGCTCC TGGGGAGCAG GATAAGACTT GGTACTTACC AAAGCTCCCG
167281    GGCCCACACA CTCACCTTGT AGCCCTGGCA TACGTCTTCA ACAAGAGCTG TGGTGTGCCC
167341    TTTGTGCTGT GGTGCCCGCT CACAGCGCCA GCAGATGAGC TGCCCCTCGT CTTCGCAGAA
167401    CAGGTGGAAC TGCTCTCCGT GTTCCTCACA TGACATTTCT TGATCCGTCT CTTTGAGGGC
167461    TTCAATGAGG CTTCCCAGCT GCTTGTTGGG TCGGAGGCTA TCCATATGAA ATGGAGCCCG
167521    ACACTGGGGA CAGCAGAATG TCTCCTGCCT CAGTTGCTTT TGGCTTGGGT TTTTAAAGAA
167581    GTCTGTTATA CACAAGTGGC AGTAGCTGTG TCCACAGTTG ATGCTTACTG GGTTCGTCAT
167641    CAGGCTCAGG CAGATGGAGC AGGTGGCTTC CTCCATCATC TTCTTGGTGC TGGTGGTTGA
167701    GGCCATAGCT TTTATTGAAA AGCTCCAATA TTGGCTCTAG AGATGGAGAT GAAGCAGCCA
167761    GAATTTTCCA CCGTGATGAA AATACACCTC ACCTGCACCT CTATGTGATG AGCTGGCTGC
167821    AACTGACTTC CATAGGTCTT GAAGGTTTTC CTTCCAACCC CTATTATCTC ATTTTGTATT
167881    GAAGAAAAGA GGACCTAAAA GGAAGAAGTT GAGGCTGAGG TTGTTTGGGC CACGTTTGAG
167941    AACTGCAACC CAAGTGCAGA GTTTCAAGTT GCCCTCATTA GCAAGCAGTT ACAAGTGGTT
168001    GTTTAGAGGA AAAAAGCAG TTTTAAAGCA GTTTTAAAGT TGTTTGCCAA GAATTTACAT
168061    TAAAATAGCA TAAGCTTTTG ACTGGCTATA CATTGTTCTT TGTATTACAA ATCTCGGGAA
168121    TATGTAGGTA ATAGATGAGG CAGCCAGTCA GGAACAAAAT GCTTTTAAAC ATGGGGTCTT
168181    AACTGAAGAC CTATACTCCT GCCTCACTTG TCCTGATAAA TTTTGCATAC CTCACATAGC
168241    TCAGACTGCT CTAAATTATT TCATTATTTT TCTTTTCTCA GTCTTCTAAC TTTTTTTTTT
168301    TTTTTTAATG AGACGGAGTC TCACTCTGTC ACCCAGGCTG GAGTGCAGTG ACGCTATCTC
168361    GGCTCACTGC ACCTCCGCCT CCCGGGTTCA AGCGATTCTC CTGCCTCAGC CTCCCGAGTA
168421    GTAGCTGGGT CTACAGGTGT GCACCACTAC GCCCAGCTAA TTTTTGTATT TTTAGTAGAG
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168481   ATGGGGTTTC ACCATGTTGG TTGGCTCGAT CTCTTGACCT TGTGATCCAC CCGCCTCAGC
168541   CTCCCAAAGT GCCAGGATTA CAGGCATGAG CCACCGTGCC CAGCCTCTTT TTCTTTTCTT
168601   ATAAGACAAG TTCTCGCTCT CTTGCCCAGG CTGTAGTGGA GGGCAGTGGC ATGACCACAG
168661   CTCACTGCAG CCTCGACCTC CTGGGTTTAA GCAATCCTCC TGCCTCACCC TGGCAGAGTG
168721   GCTGGGACTA CAGGTATGTG CCACCATGTC CAGCTAAAGT CTTCTCTCCA GAAAGAAGAA
168781   ATGCATTGGA ATTTAGAGGA TACACAAACA TCTAGCTGTA TAGCTAATAC AGTAGCCACT
168841   ATCATGAGTA GGAATTTAAA TTTAACTTAA TAAAAATTAA AATGAAAAAA TTCAGTTTTT
168901   CTGTTCCAGT TGCCACATTT TGATTGCTTA ATAGTTGCAT GTGACTAGTG CTACATAAC
168961   AGCCTCAATA TACAACATTC TGTTATCACA GAAAGTTACC TTGGACCAAG TGCTGGGAGA
169021   AGCAATGCAG GCTTCCTCAC AAAAGCTGTA AAGAGAGAA CTCAGGGAGT GTGAAACTCT
169081   TTCCTATTCT AGTTAACTTC AAGAATAATT GTTACCAGGC CAGCACGGTG GCTCACGCCT
169141   GTAATCCTAG CACTTTGGGA AGCCGAGGCG GGCAGATCAC CTGAGGTCAG GAGTTTGAGA
169201   CCAGCCTGAC CAACATGGCA AAACCTCATC TCTACTAAAA ATACAAAAAG TTAGCTAGAT
169261   GTGGTGGTGC ACACCTGTAA TCCCAGCTGC TCAGGAGGCT GAGGAAGGAG AATGACTTGA
169321   GCTCCGGAGG GGGAGGTTGC AGTGAGCCCA GATTACACCA CTGCACTCCA GCCTGGGTGA
169381   AAGAGCGAGA ATCTGTCTTA AAAAAAAAAA AAGAATAAT TGGTACCAGA ATTACTCTTT
169441   GTAATTAGTA GTAACACTTA TGCAATTGGG TGATCTGTGA CAGATTCCAT TGAAGGAGTA
169501   TGGGGAGCTT CACCCCAATA TATGACTCCC TGGTATAATG AGTATTTTGA ATTAAAGGCC
169561   CTTAGAGATC AGCAGATGCT GGAAGAGACT TTTCCCCTAT CTACATAAAG ACCAGTCACA
169621   CTAGACAAGA AGAACAATTG TTTTTCCTTC CAACCCCTAT TATCTCATTT TGTACTGAAG
169681   AAAAGAGGAC TAAGAATGTA ACCAGACCTA ATCAGACACT TTCACAAAAT AATGTCTGTC
169741   TCTCAGGCTC ATTCATTTTC CAAAGAGAAC CATTTACAAG TTAAACTCTG TTCCTCCATT
169801   CATTCATCCT CCCAAATATT CATTTATTCT CCCTAGTAAT CATTTACTGC CCCTCAAAGA
169861   ATTACCTATA TTCTCCTGAT ATCACCCTTC CCCTCTGAAA TAAATATGTA TACATGTATA
169921   AACGTTATAC ATACATATTT ATACAGTATA CATACATATT TATACATACA TACATATGCA
169981   TACATATTTA TATTTATGTA TTTATACATA AGTATTTATA AATAAGGCTA TATAAGTATC
170041   TACCCCCATT GGCAGAGGGG GTAATCACTC TGTGATTCTA GCCCATGTAC TTGTTAATAA
170101   ATTTGTATGC CTTTTCTCCA ATTAGCCTGC CTTTTGTGAG TCGATTTTTC AGTGAACTTC
170161   AGAAGGCAAA GGGGAAGTGT TCCCTTGGCT CCTACACCAT CATGACAATA AAATTTGACT
170221   CCACCTCGAC CCCCCCCATC CCCCACAAAG AACAACAACC AACACTGGTT AATAAGGTCG
170281   GTTGTTTTTT GTTTGTGTTT TTGTTGTTGT TGTTTTTGCT TTCAGGAGCA GAGGTATAAT
170341   AGGCAAAAGA AAGAGAAAGG AGAATAGTGA ATACCTCTTC TGCAGAGAGG GGTGCCTAAG
170401   TGGGACTTCC CTGGCTAATA ACGTCTTGCT AGAGACCCAA CCAGGAGGAT AATGGAAGCA
170461   ATCAAGGCAA CCAGAACAAC CAGAAGAACC GGTTTATCCT TTTTGTGCCC TCTCCCTAAA
170521   CTGAGGGAAT AAGAATTGGA AAGAAGGCTG CAGAGCAGAG GGTTTGCTCC TGAGGAGCAG
170581   TTATTTCTAT GGGATCAGAG CTCCTGCAGA ACTGGGGAGT TTACTTTTAC TATCTCTTCT
170641   CCAGGACAGG ACCTATCTCA AGAGACATGT TCAGAGTGAT TGCAACATAA AGAGTTTGCA
170701   GACCCAAGGA GGTAGGGAAG GCAGAAAGAA GATGGGGGAG GCCAGGGATA GGCAACAGAG
170761   GAGTGACCAG GAGCGAAAAA GCCTGCCTCT TCTGAGAACC TAGCTGGGCT CTCCCTGTAC
170821   CCCCGATCCC TCCCCCCCGC CCGCCCCAC ACCCCTACTC CTGGGAGCTC CTCTAGGACA
170881   GGGGCAGAGT CAGGAGGAAG TTTGAAGAGT GCCTAGAATA AAAAACAGTA ATTTAACTAC
170941   AATTACCGGG TAGGCTGTTT TCCTCTCACA ATTTGATCAG TCTCTTGAAG CCACACAGAA
171001   TTTCTTCTGA AGACGTGTAT TCCTTGGCAG GCTATTTCCT CCAGTGATAC ACCAGGCCCC
171061   TCTCTGCTGG GGTCACTGCT CTTCTGGGGA GATGGGGCTC CCCTCCTTCC AAGGCTCCAG
171121   GGTTCCTGTC CTGGGCCCCA CTCATCTAAG TTCTGAATCT TCTGAGATTT GGTGTAAAGT
171181   CTGGTGAAAG AAAGAGCAGG AAAGAGGTGA GAGCTGTAAA ACAAAGAAAG TCCTGACCAT
171241   TTTCAGAGTT GGAGGGGCCC TGCTGTCACG AAATATATTC CCCACCCCAC TTGCCATCAG
171301   TACACACTCA CATATCCACT GAGAAAACCT TAGCCTGGAC CTTTTCCGTA ACCTTCACTG
171361   CTCAGACACT TACATATTCG CTGCTAGTCC CCTCTGTTGC TGCCACTTCC TGGGTCAGGA
171421   AGTTAACTCA GACCGGATTA AACTGAGAAG TGAAACTACT GTGGGAGGCG GGGCTCATAA
171481   GATTTAGGAG AAAACTAGTG ACGTTGTTCA TATCATTTGC ACTCCGCCTC TCCGGTAAAG
171541   GAGGGGGAAA CGTAGGAAGA AAATATCCTT CTTTTACAGC AATAAAAAGA AGGAACCAAT
171601   TAATAACCCT GTAAACTATC ATGTGACCCC AACACAGAGT ATCTAAAAAC AGGAAGCCTG
171661   CAGAGGTTCA GTTCACAGAC TCTGATTTGA GATCTTTCTA CTTTTGCCAC CAACTCCCTT
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171721   GGGAGTCCTT AAGCCTTCCT AGCTGATGTT ACTTCTTTTG CTATTTATGG GTTGCTTGTG
171781   GTTCTATAAC TGCTCTGAAG GGTGTGGTGG AAAAAGGGGT GGTAACAGCA GTAGGACTCA
171841   TTGGCATCAC AAAATTCATC TGAGTCAGCT TTCTATTCTT CTCTGTCCCG TTCTGTGTCT
171901   TGTTTTTCTC CTTGCTGTCC TTCTGCAGGA CTCAGATCTT CTTCAATAGC GAGGGTCAGC
171961   CAGGATAGAA AATGGGAGTC ACTAGTGGCC CAGCAGTGAG TGCCCCAGC TTAGAGCTGT
172021   GTGGGATCCC TGGGACCATC ACTCTGCTTT GTGCTTTGTG GAGAAAAGGC TGTGGGGTCC
172081   AGGGTCAAGT CCTTAATGAC TTAGCTCCAG CTTCTCCACT TCAAAATGAA AGGAAAAGTA
172141   CTATCACCAC CCGTTAGAAT TATTATTTCA TGGGGAAAAA AGATGGATTA CTATCTCACA
172201   ATAAGAGCTT GTCACATTTA TAAGTCTCAG GTGTAAGAGG CATTTATGAT AACAACATAA
172261   TAAATGCTGG CTTAAGTAGA TGCAGTGGTC CAAGGGAACC AGTAAGGGGA GCTCAGGACA
172321   CAGGTGGGAG GAGAAATTAA ACTTGAATTC TGGGAGCCAC TGGCCTGTCT GGGCCCCTGG
172381   CCTGCCTGCT GACCCTGATA GCCAATGGAA CATGGAGTTT GGCCCAGCTG CAATCCCTCT
172441   GGTCCAACTA CTCAAAATAA AGGCAAGATT GGGAAACACG TTCCTTTCTT CCTATACCAA
172501   GCAGAAGACT CTTCAGCACT GCACCCTCCT GGGTGCTCAC AGAGCCTTCT GTTGTTTTGC
172561   CACCTACGAT TCATCATGCC CTGGCATGAT GGTTGCAGAC CCCATGCATA GCATGGGACA
172621   TTCTACTCCT GAGGCAACCA GCACACAGAG AGAGGAGAAA GAATGAGCCC CTGAATCCTT
172681   GGTCCCACGA TGAGTCCTTG CAGATATCTA CAACTTTCAT TGTTGTGGAT GTGACTCTGT
172741   ACCCAGGCAT GGCTCATTCC AGATCTGTCC TATTGTCAGA GGTGTTCAAA CCAGAATGAC
172801   TCCATTTTGA ATGGGGGCTA GGTAAAATAA GGCTGAGACC TACTGGGCTG .CATTCCCAGG
172861   AAGTTAGGCA TTGTAAGTCA CAGGATGAAA TAGGCAGTTG GCACAAGACA CAGGTCATAA
172921   AGATCTTGCT GATAAAACAG GTTGCAGTAA AGAAGCTGAC CAAAACCCAC CAAAATCAAG
172981   ATGGCAACAA GAGTGGCCTC TAGTCATTCT CATTGCTCAT TATACACGAA TTATAATGTG
173041   TTAGCAAGTT AGAAGGCATT CCCACCAGCT CCATAGTGGT TTATAAATAC CATGGCGATG
173101   TCAGGAAGCT ACCCTATATA GTCTAAAAG GGGAGGAACG CTTGGTTCTG GGAATTGCCC
173161   ACATCTTTCC CAGAAAACAT ATGAATAATC CACTCCTTGT TTAGTACATA ATCAAGAAAT
173221   AACTGTAAGT ATCTGTATTA GTCCATTTTC ACACTGCTGA TCCAGACATA CCTGAGACTG
173281   AGTAATTTAT ACCAGGAAAA AATGTTTCAT GCTCTTACAG TCCCACGTGT CTGGGGAGAC
173341   CTCACAACCA CAGCAGAAGG CAAGGAGGAG CAAGTCAGGT CTTACATGGA TGGCAGCAGG
173411   CAAAGAGCTT GTGCAGGGAA ATTCCTTTCT ATAAAACCAT CAGGTCTCAT GAAACTTATT
173461   GACTATCATG AGAACAGCAG TATAAATTAC TCAGGGAAAG ACCTGCCCCC ATGATTCAAT
173521   TACCTCCCAC CAGGTCCCTC CCACAATATG TGGGAATTTA AGATGAGAGT TAGGTGGGGA
173581   CACAGCCAAA CCATATCAGT ATCCTTAGTC CAGAAGCTGA TGCTCTGCCT GTAGAGTAGC
173641   CGTTCTTTTA TTCCTTTACT TTCTTGCTTT CACTTTACTG TGTAGACTTG CCCCAAATTC
173701   TTTCTCACAC GAGATCTAAG AACCTTCTCT TAGGGTCTGG GTTGGGACCC CCTTTCTGGT
173761   AACACTATCA AAGGATCAGG AAAAGGAAGC TAGTGAATGC TAAAAAGGAA ACAAACTACC
173821   ATTACCAATA ATAACAGCAA GACAAAAGCA AAACGGATTG TGACAGCTGT CCCATCTCAC
173881   ACCTGTTTCC CATTGCAGGA AGGAGGGGCT GGTTCATGCA CAGAGTGGCC AATATTAGAA
173941   GCAGAGATGG GGTGCAGATG AGACTTCAGG AATATGTTGA CAAAGGCAGG CCTAGGGAGA
174001   AATCAACCTG AACTATCCCC AAGGAGGAAT GCATTATCTC TAATATGTAA AGTTAGGCTT ·
174061   GATCCTGTGA TTATGGGATA TAGGAGTCCA AAGACTCACA ATGGGAAGTA GGTCACTAGA
174121   GTCTCCTTCA GAAGCTCTGT ACTGTGTGTT CCCACTGTGG GCAAGAGTCA GCACTCAGCT
174181   ATTCCTAGAA TGCCTTTCCT CAACTCCTTC AGATTTTGCC TCTCAACTAA CCCTATCCTG
174241   ACCACTTGTT AGCAAGTGTA CCCCTCTCTC CCTCCCAAAC ATTTTCAAAT CTATTTTGTT
174301   CCCATGGCAC TTATCACTGA ATATTTTACT AATTTATTTT GTTTAGTGTT TGCTTCCCTC
174361   ATGAGAATGC AAAGGGATGG ATTTTTTTCA ATATTGTTCA CTGATGAATC CCAGTAACTA
174421   GAATATTTCT AAGCATAGTG ATGTGCATTA AATCAAAGAG TAACTTTCTG AATTGCACTA
174481   AACACACATC ACAAGAGGTG TGTGCACATA TGTGCATGAT GCACGTAGTG TGGTGTGGGT
174541   GTTGTGTGGG GTATGTGGTA CTGTGTGTGC TGTGTGTGGT ATGTGATACA TAGTTTGTGT
174601   TAGTGTGATG CATGTGATGT GGTATGTGTG TGCGTGTCCA TACATATTAG GGGTGGCGGG
174661   GATGTTAATA TGTCAAATGG TACTAGAAAG TATCAGAACT CATGGTGCTT ACTGGTTTCC
174721   CAGAGAGCTG CTTCTCTCCC ACCTGTAGGA TATACTGATG GTTTGGACAG AGAAGAAATA
174781   AAAAGAAGGC TGTGACCTAC TGGGCTGAGG AAATAAAAAC GAAGTAAAA GAAGAGCTGG
174841   GAAAAGAGAG TGGAGGGGCC AAGGGAAATT TCCCCTTTGG CTTCTGGGGA AACTTTGCTG
174901   AAAAATCAAC TCACAAATTT ATTAACATGT ACACAGGGAG AACCATAGAA TGATTATCCA
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174961   CTTCCCAAGA GGGCTTAAAA GCTTATATAT TATCCTGGCA AAACAGATTA TGGGAGGGGA
175021   AGAAGAGAAA CTCTGTTGAT GGGATTACTG TTGCGGATTT TTGCTCCTTC GCTCAGCTAG
175081   GTCCGGGTTT TTGTCTCACA GCCAGGAAGA ATTAGGCATG CAGCCATCAA AGAATGAGTG
175141   GAGTAGAATT TATTAAGTGA AAGGAAAGCT CTCAGCAAAG ACAAGGGTCC TGAAAGCAGA
175201   TTTCTGGTTT GCTCTTCACA GTTGAATACT AGGGCTTAAG ACTCAAATTC CTGACAACTC
175261   CACCCTGTCC TACCAGTGCA TGCAGGCCTT TAGACTGAGC TACTCCATAT TGATTAATTT
175321   CCTGAACTGT GCATGTGTTA AGGAAAGGAA TCATCCACTG CAGGCATGTT TAGGCAAGCC
175381   CCCTGTGCAA GTTCCCTTAT CTGCACAAAA CATCCGGTGT AAGCACTTGT GGGGCAGGTC
175441   AGAGGTTCTC TGGGTACCAT TCCCTTACTG TCTGCCTAAA GCAAGCTGGC CAACTCCTTT
175501   CATTACTAGG GAGAGTAAGT AGATCAGGGA ACAGAGATTA ACTTGAACAT TATCTTGTGA
175561   AAGTCCGTTC GGGCATGGTT ACATTCTTGG TCTTACAGGA AGGGTAAATA AAAATAATTG
175621   CTCTTTTTGG TGGGTCTGGA TCTTAGGTAG ATAAAGAAAC TTTAATTCCA CGATGTGTTT
175681   TGGTAGGGAT AGTTGGTGGC AGGGATGTCA GAGAGACTTT GAGGCTTCTT CAGTTCAATA
175741   TGACCAAGGG CCATATATTA GGGTATCAAT TTCTGAGCCC CAACAAGAGC TTAGGAGAGA
175801   TGTGATAGCA TCACAGTGTG AAAGCAATTT TTTGTTTGTT TTTAGAGACA GGCTCTTGCA
175861   CTGTCACCCT GGCTGAAGTA CAATGGTACG ATCACAGCTC ACTGTAATCT TGAACTGGGT
175921   TCAAATGATC CTCCCATCTA AGCATTTCAA AGTGTTGGGA TTACAGGCAT GAGCCACGGT
175981   ACCCAGCCTG AAACTGCACC CACTTTCTGA TAAACTTTTC AAATGACTAA AGGGGAGAGA
176041   GTAAGCACTA CTCAGAGGTA GGAAGAAAGG ACACAGGATT ATAGGATTAA AACAACAACC
176101   ACCAAAAAAA ACCAGACCGG TGTGGTGGCT CACACCTGTA ATCACAGCAC TTGGGGAGGC
176161   TGAGGTGGGG GGAGTCACTG GAGGCCAGGA GTTCGAGACG AGCCTGGCCA ACATAGCAAG
176221   ATGCTGTCTC TATTAAAAAA AAAAAATACC TGCCTTGAGC TAATCAGAAT CATGGACCCT
176281   GACAAAGGAT GTCCCAAAGT AAGTCTTAGC ATTTTTTTTT TTTTTTGAG ACAGTCTCGC
176341   TGTGTTGCCC AGGCTGAAGT TCAGTGGCGT GATCTCGGCT CACTGCAACA GCTGCCTCCC
176401   AGGCTCAAGC AATTCTCCCT GCCTTCAGCC TCCCAAGTAG CTGGGATTAC AGATGCCCAC
176461   CACCACGCCT GGCTAATTTT TGTTTTTTTT AATAGAGATG GGGTTTTGCC ATGTTAACCA
176521   GGCAGGTCTT GAACTCCTGA CCTCAAGTGA TCTGCCCACC TTGGCCCCTC CATAGTGCTG
176581   GGATTACAGG CGTGAGTCAC TGCACCCGGC AAAGTCTTAG CATTCTTTAC AAACAGTTTG
176641   TACCCGTATC TCTAAAAGGG AGTAGTGAAT TTCACCCCAA AATGTGGCTT CCTGATATAA
176701   TGAGTATTTT GAATGAAAAA CTCTTAGAGA TCAACAGACA CTAAAGAGAC TTTTCCCTAG
176761   GTACATAAAA ATAGGATGGC CCCACCAGCG AGAACAATTG TTCTTTTCTC CCTCTCTGTT
176821   ATCTCATTGT GCATTATAGG AAAGACCAAG AATGTAACCA CACCTGAACA GACCCTTTTA
176881   TAAGATAATC AGTCTCTAAG CATCATTTAA ATTCCAAGGA GAACTATTTA CAAATTTATC
176941   TGTTCTTTGA TCCAATTAGT CTCTCCTGGT AGTTACATAT TGCCCCTCAA CAGAATTCCT
177001   CTTCTTCTGT TTCCCATAAC CTATTTTGCA AGGATCAAGC CCCTGTTATT TCTTCAACTT
177061   CAAGGTGGCA TATAAGCTTC TAAATTCCAC TGGGATATTG GTACTATGTG CATGAGGAGA
177121   ACCACAGAGT AATTAAATTG TAAAGCCTTT TATCTTATGA ATCTGCCTTT TTTTGTGTTC
177181   ATTTTTCAGC AAAACTTCCA AGGGCAAAGG TATAAAACAA AAATAAAATT CTAAAGCCCC
177241   CCAACCATCT GAATAGACTT TCTCTTCAGT CAGGCTTCTT AAAATGTAAC CTGAAAGACT
177301   GGCTCAGGCC ATTAAGGGAA GTGGGGGTTG AACATGCCTC ATTATTCCTC TCTGGCATTA
177361   ACATCAACAC AGCTTTTAAG TCTGATAAGA AACATTTTAC AACCTATTCT CTCTGAAGCC
177421   TGCTAGCTAA AAACTTCATC CCATAGTACA ACTTTGGTCT TCACAACCTG TTATCACAAC
177481   CTAGTGCTCC TTTCTATTAA TCCCAAATCT TTATACAAAC TCAACCAATT GTCATCACCT
177541   CCACCCCACT CCTCCGCTGC TTCCAGTTGT CCCGCCTCTC TGGACCAAAC CAGTGTACAT
177601   TTCTTAAACG TATTTGATTG ATGTCCCATG CCTCCCTAAA ATGTATAAAG CCAAGGTGCA
177661   TCCCAACCAC CTTGAGCGCT TGTTCTCAGG ACCTCCTGAG GGCTGTGTCA TGGGCCATGG
177721   TCACTCAAAT TTGGCTCAGA ATAAATCTCT TCAAATGTTT TACAGAGTTT GGCTCTTGTC
177781   ATGACACAGA TGACTGCTTC ACTGAAGCCT GCTCTGGAAG TGAGTGGGGG TTTTGCAAGG
177841   ATAATTTTCC CCGGATAGCC CCAGAAGCAG CTAGTAATAA TACACTTAAA GGTAGCTAAA
177901   ATGCATTGAA CACTTGTTTT GTGCCAGACC TATGTCAACA TTTGCTTTGT GCCAGGCTTA
177961   TGCCAGTACT CCTGATTTGT TAATACATTC TAAATAAAAA TTCTGGAGTT TCAAATATAA
178021   TAACTGAAAA ACAGAAAATA AATAAAAATA TATAATAACT GAAATAAAAA TTTACTAAGG
178081   CTGGGGATGG TGGCTCACTC ACACCTGTAA TCCTGTTACC GGAAAGGGGT CCGTCCAGAT
178141   CCAGACCCCA AGAGAGGGTT CTTGGATCTC ACACAAGAAA GAATTCGGGC GAGTCTGTAA
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178201  AGTGAAAGCA AGTTTATTAA GAAAGTAGAG GAATAAAAGA ACGGCTACTC CATAGGCAGA
178261  GCAGCTCTGA GGGCTGCTGG TCGCTCATTT TTATGGTTAT TTCTTGATTA TGTGCTAAAC
178321  AAGGGGTGGA TAATTCATGC CTCCATTTTT TAGACCATAT AAAGTAACTT CCTGACGTTG
178381  CCATGGCATT CGTAAACTGT CGTGGCGCTG GTATGAGCAT AGCAGTGAGG ACGACCAGAG
178441  GTCACTCTCA TCGCCATCTT GGATTGGTG GGGAGCAGTG AGGATGACCA GAGGTCACTC
178501  TCATCGCCAT CTTGGATTTG GTGGGGTTTA GCCAGCTTCT TTACTTTTTT CTTTTTTTTT
178561  TTTGCCCAGG CTGGAGTGCA GTGGCACGAT CTCAGCTCAC TGAAACCTCC AATTTCTGAG
178621  TTCAAGCGAT TCTCGTGCCT CAGCCTCCCA AGTAGCTGGG ATTACAGGCA TGTGCCACCA
178681  CACCCAGCTA ATTTTTTATA TTTTTAATAG AGACCGGGTT TCGCCATGTT GCCTACGCTG
178741  ATCTCCAACT CCTGCGCTCA AGCCATCCAG CCACCTTAGC CTCCCAAAGT GCTGGGCTTA
178801  TAGGTGTGAG CCACCCCACC TGGCCTAGCC GGCTTCTTTA CTGCAACCTG TTTTATCAGC
178861  AAGGTCTTTA TGACCTGTAT TTTGTGCCCA CTGCCTGCCT CATCCTGTGG CTTACAATGC
178921  CTAACTTACA GGGAATGCAG CCCAGCAGGA CTCAGCCTTA TTTCACCCAG CTCCTATTCA
178981  AGATGGAGTC TTTCTTGTTC AAATACCTCT GACAAGCCCA ACACTTTGGG AGGATGACAC
179041  AGGAGGATTG CTTTAGCCTA GGAGCTCAAG ACCAGCCTGG GCAACACAGT GAGACCCCAT
179101  CTCTAAAAAA AAAAATACAA AAAAATTAGC CAGGCATGAT GGTGTGTGCC TGTAGTCCCT
179161  GCTACTCAGG AGGCTGAAGT GGGAAGATGG CTTCAGCCCA GGAATTCAAG CTGCATTGT
179221  CAGAGGCATT TGAACCAGAA TGACTCTATC TTGAATAGGC GCTGGATAAA ATAAGGCTGA
179281  CACCTGCTAG GCTGCATTTC CAGTATGTTA GGCATTCTTA GTCACAGGAT GAGATAGGAA
179341  GTCAGCACAA GGTACACATC ACAAAGACCT TGCTGATAAA ATAGGTTGTG GTAAAGAAGT
179401  TGGCCAAAAC CCATCAAAAC CAACATGGCC ACCAAAGGGA CCTCTGGTTG TCTTCACTGC
179461  TCATTATATG TTAATTATAA TGTATTAACA TGCTAAAAGA CACTCCTACC AGCATCATGA
179521  CAGCTTACAA ATACTGCGGC AATATCTGGA CTTTACCTTA TATGGTCTAA AAGGTGGAGG
179581  AACCCTCAAT TTTGGGAATT GTCCACCCCT TTTTTGGAAT GCTCATGAAT AATCCACCCC
179641  TTGTTTAGCA CATAATCCAG AAATAACTAT AAGTATGCTT ATTTGAGCAG ACCACGCTGC
179701  TGTTCTGCCT ACAGAGTAGC CATTCTTTTA TTTCCTTACT TTCTTAATAA ACCTGCTTTC
179761  ACTTTACTGT ATGGACTTGC CCTAAATTCT TTCTTGTGTG AGATCCAAGA ACCCTCTCTT
179821  GGGGTCTGGA TCAAGACCCC TTTCTGGTAA CATCTTTCTG GTGACCACGA AGGGACAATA
179881  CTGAGGAGAC TCTGAAGCCA AAGGAAACAG ACTACAGCAC CAACTGGCTG ACTTTGGGTA
179941  AGTGGTGGAG TCCCCGGGTA AAGGATAGGA TTGGGTTAGA GGTGCAACTT AGGGGAGATA
180001  GGGTCTCTCC TAAGACAGAG AGGGTTTCAG TCCGCTCTTA ATAAAGGGCA AGAATGCTTG
180061  ACCGAACTTG GGTTTGAGAC CCAACTTAGG AAGGCTACAG TCCTTAAGAT TTAAGGGGTT
180121  AGAGGCCCCT CTCAGTAAAG TCTCTCTTGG TTAAAAACGG ATTTAGCATT AGGGGATGTT
180181  AACTGCTATT CTGTTTGTAT TAATCTTCCC TGTGCTCTTT GCTGACAGCT ATGGGTGACA
180241  GGATTAGGCA TGTACAGGAT CACGGGACAT TGGGAACTTT TCTTCTCTCC AAAAGGGGAA
180301  GCTTGACAGC TGATAGGACT GTTGGAAAAG ATCCCTTTGC TATGACAAGC AGCCGCCTGA
180361  ACTTTTGATT CAGTGTTGCT GCAATGGGTG GGTCTTTCTC TGGCCTCTGT GAACTCCTCA
180421  CCTTCCCCAT CTCACCACAG GCAATGCTTT TCTCCCTTTC TCTCTTTTCT CTTTTCTGTC
180481  TTTTCTGTTA CTTGAGACAA CCATCTTGCC CAGAGACCAT ATGTTGAAAC TCCTGGTCAG
180541  AAGTTTGATT AAAGATGAAA GGGCCTATCT GGGGGCAAGT TTGAGCCTTC CCAGTTAGAT
180601  ATTGGGTGCT AAGTGGAGTG GCCAATGTCT ATGTTTTGTC ACATGTATAT TGCTCTGGCT
180661  GAAATGGAAA ACGTTAATTT GGTTACTTTA TGTGGCCATT GGGCAGCATC TTACAAAAGT
180721  GAGAGACATT TATTTGCCTG TGGTTCCATG AAACAGAAAA AAGTTGGTTT CTTTTTGTGT
180781  CGTAGCTTGG ACCCAAGGGC TTTGCAGTGA GCAAGGTTGC TAGTGCTGCT CAGTGAAAGA
180841  GAACCCAGAA ACCTGGCATG CCAGCAAAAG GGTAAAGATT CTTACCAGT CAGGCTTCTG
180901  GCCTCTCTCT CTTAGTGAAA ACTGAATGAA TGGTAAAAAT CACTGTTTAT CACCTCTGTA
180961  AAGTTTTGAT TAATGGGAAC AAGGATTTGT GGGGCTAGTC TTAAGCTGTA ATGAATCTGG
181021  TATACTTTGT GATATCAATT TGTCTTTCTG TATTACTCTG TCATAAAGAG GAATATGGTA
181081  GGATAGAACA TGGGCTCAGG ACTCCATAAG CCTGCTGTTC AAGCCAGCCC AGTAAACTGG
181141  TCCGTTGCAA AGTTTATTAC AGGTCCCTGG AAAAAAAAAA AAATAAAAAC TGGATGAAGT
181201  TTCCTTCTCA TCTTGTTTTA TGTCCTTTGG AGCTTCACCT TGTAACCACG TGGCGGTACT
181261  TTCTCTTGGT CTCTGCCATC CAGGGAACAG GAATTTTGGG GTTTATGTAA TAGTTAACTC
181321  TAAAAATTAT CTCAAGCCAT TGCAAGCTCA AAATTGGCTG CTCTGGACCC CTTCTGGGAA
181381  GGGCAATGGA AACTAACCAG TGTTGTAGCT CAGCAGCTAA GGATTTGTCA TTTTATAATG
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181441   GCGGCCAAGG TTCAATCCTG GCTTAGGGAA TGAGTACTTT CTGATTGATA TCTGTGTGAC
181501   CTTTACCATT TGTTGATTCT GTTCTCTTCC CCTCCACACA CTGTCTTGAG TTTTCCTCTC
181561   TCTGAGAACC TGGGAGATTA TCTTTGGTAA AGTTCAAAAG CCAGAAATAA TGGCCGTGTG
181621   GGATGGCTAA AGTTGAGTAA TAAGAAACTT AAAAGGACTC CTTTTTTTTT TGCTTTAGAG
181681   TGCTATGGTT TATGGTTAAA AGCTTAATTA AAAGTGGATA TTCAATCTCT AAAAGCCTGG
181741   GACTCCTTGG GAAAAGCAGA GGAGGCACCA CAGACCCCAT TTTGGGAAAA CCTCTGTTTT
181801   CCTCATGAAA CCCCAGGAAC TGGAAGTGGA TAGATCCTTC GCAAAATCTA AGGCTCTGTT
181861   TGGCTTTGCA TTATGTTATC TGATGTTTTT GACTTTGGGG GGTATCAGAA ATTACTTTGC
181921   ATTATGAGGG AGATCTGGTG TGTAATAACC AGGTAGGAAA TATACTTCTG GGGATAGCTA
181981   AAGGCAAATA TAGGTGAATA CTTGGCTATT TGCACTTTTG GATCACAAGA AGCATTCTCT
182041   TGACTACCTA GAAGGTATGG AAATGTCTCC ATCCCCACCG AGAGATAAGA TTCCCAGGGG
182101   AGATGGCTGA TCCCCCAAAA GAGGGCTGAT TCCCTCTTTT GGGATCCAGG ATCTGGTATA
182161   AAAATGGGAC CCTGGCCAGG CACAGTGGCT CACGCCTGTA ATCTCAACAC TTTGGGAAGC
182221   CTCAGAGTTA TGAATGTCTC ACCATACTGA CACTTTGTGA CTGAGCTCCT CTCTACCCTG
182281   GACACAAGAG ACCCTAATAA TTAGACAGGA ATATCATTGC CCCTATTTAG TCTGAAGAAG
182341   TTATAGAAGA CGGATCTTTA TCCCACTGCA ATCCTTAGGA TTAAGGGTTC CCTGGTAAAA
182401   GGGAGTGGGA AAATATGTCA GAGGCATTTG AATCAGAGTG ACTCCATCTT GAATAGGGGC
182461   TGGGTAAAAT AAGGCTGAGG CCTGCTGGGT TAGGTTAGGC ATTCTAACCA GGAGTTTAGT
182521   CACAGGATGA GATAGAAGGT TGCACAAGGT ACCCGTCACA AAGACCTTGC TGATAAAATA
182581   GGTAACGGTA AAGAAGCCAG CTAAAGCCCA CCAAAACCAA CATGGCCACA AAAGTGACCT
182641   CTTGTCATCC TCACTGCTCA TATACACTAA TTATACTGCA TTAGCATGCT ACAAGACACT
182701   CCCACCAGTG CCACGACAGT TTACAAATAC CATGACAACA TCTGGACGTT ACCTTATATG
182761   GTCTAAAACG GGGAAGAACC CTTAGTTCTG GGAATTGTCC ACCTCTTTCC TGAAAAATTC
182821   TTGAATAATC CATTAGTTTA GCACATAATC CAGAAATAAC TATACGTCTG CTTATTTGAG
182881   CAGTCCATAC TGCTGCTCTG CCTATGGAGT AGCCATTCTT TTCTTTTATT TTTATTTTTT
182941   AGATAAAGAC TCGCTCTGTC ACTCAGGCTG GAGTCTGGAG TGCAGTGACG TGTTTTGGCT
183001   CACTGCAACC TTCACCTCCC GGGTTCAAGC AATTCTCCTG CCTCAGCCTC CCAACTAGCT
183061   GGGACCACAG GTGGGTGCCA CCATGCCTGG CTAATTTTTG TATTATTAGT AGAGATGGGG
183121   TTTCGCCATG TTGGCCAGGC TGGTCTCGAA CTCCTGGCCT CAAGCGATCC ACTTGCCTTG
183181   GCCTCCCAAA GTGCTAGGAT TACAGGCATT ACCCACTATG CATGACCCAT TCTTTTATTT
183241   CTTAACTTTT TTTTGTTTTT TTGAGACAGA GTCTCACTCT GTCACCCAGG CTAGAGGCTG
183301   GAGTGCAGTG GTGCGATCTT GGTTCACTGC AACCTCTGCC TCCTGGGTTC AAGCGATTCT
183361   TCTGCCTCAG TCTCCTGAGG AGCTGGGACT ACAGACATGT GCCACTACAC CCAGCTAATT
183421   TTGTATTTTT AGTAGAGACA GTGTCTTGCC ATGTTTGTCA GGCTTGTCTC GAACTCCTAA
183481   CCTCAAGTGG TCTGCCTGCC TCAGCCTCCC AAAGTGCTGT GATTACAGGC ATAAATCACT
183541   GCGCTCGGCC CTTCTTTACT TTCTTAATAA ACTTGTTTTC ACTTTACTGT ATGGACTAGC
183601   CCCAAATTCC TTCTTGTGTG AGATCCAATA ACCCTTTTGT GTGTGAAAGA ATGTATTGCT
183661   GCTGTTCAGG CTGGAGCAAG CTGGAGCTCA TGCTGCTGCT CAGACTGGAG CATGCGTGAT
183721   CTGTGATCCC AGTAAGAGGA TCATGGTCAC TCCAGCCTGA ACGACAGCAT GATATCTCAT
183781   CTGTAAGAAA AAAAAATTAC TAGAGGGCTT TAACAGCAAA TTTGAGCAGC AAAAAGAAGT
183841   AATCAGTGAA CTCAAAGATA GGTCAATTGA AATGATCTAC TCTGAAAAAC AGAAAGAAGA
183901   CAGAATGAAG AAAAAGAAAT AGAGCCTTAG AGACAGGGGA TACCATCAAG CATACTAATA
183961   TATGCATAAT GGGACTCCTA GAAGGAGAAA AGTGAGAGGA CAGGGAGAGA GAATGTTTGG
184021   AGAAATAATT TCTCAAAGCT TCCCATGTTT GGCAAAAAAG CATTAACTTG CATACATATT
184081   TTAGGAGCTC AATGAATTCC AAGTAGGATA CACTCAAAGA GATCCATACC TAGACACATC
184141   ATAATCAGAT TATCAAAAGA TGAAGAAGAT GAATCTTGAG AGCAGAAAGA AAGGAACAAT
184201   TCATCACATA CAAATAGTAC TCAAAAGATG TCTGGAGTAG GTATACTAAT ATCAGACAAA
184261   ATAAACTTTA AGATAAGCAT TGTTATAATA AATAAAGAAA GGTATTTTGT AATGATAAAA
184321   GTGTCAATTC ATCAAGAAAA CATAACATTA TAAACATACA TGCACCTAAC AACAGAGCCC
184381   TAATATTCAT GAAACAAAAC TGACAGAATT GAAGGGAGAA ATAGAAAATT CGACAATAAT
184441   AGTTGGAGAC ATCAATACCT CACTAGTTAG ACAAGATCAA CAAAAAAATA GAAGACTTAA
184501   CACTTGAAAA CACCTAACCT GACCCTAACA TAAATCTATA GGTCACTACA CCCCAAAACA
184561   GCAGAATAAA CATCCTTCTG AAGCTCACAT GAAACATTTT TCAGGATAGA CTGTATATTA
184621   CTTCATGAAA TAAGTCTCAA TAAATGTAAA AGGACTATAA TAATAGAGTA TATATTCTCT
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184681    GACCAAAGTG GAATGAAGAT AGAAATCAAT AACTAGGCTG GGCGTGATGG CTCACGCCTG
184741    TAATCCCAGC ACTTTGGGAG GCCAAGGCGG ACAGATCACG AGGTCAGGAG TTTGAGACCA
184801    GCCTGACCAA CATGGTGAAA CCCTGTCTCT ACTAACAAAA TACAAAAATT AGCCAGGCCT
184861    GGTGGCATCT GCCTGTAGTC CCAGCTACTC GGGACACTGA GGCAGGAGAA TCACTTGAAC
184921    CCAGGAGGCA GAGATTGCAG TGAGCTGAGA TCGCGCCACT GCATTCCAGC CTGGGAGACA
184981    GAGCGAGACT CCGTCTCAAA ATTAAAAAAA AAAAGAAAC TAGAAAAATA AGAACAAATC
185041    AAACCCAAAG CAAGCAAGAG GAAATGAAA AATTTCAAAG CAGCCAAGAA CAAAAGGCAC
185101    ATTATGTACA GAAGAACAAG TGTATAGATC ACATATTTCT CATAGACACA ATATAAGCAA
185161    AAAGACAGTG GAGCAAAATT TTTTAGATTA ATGAAAGACC TACAATTCTG TACCAAGCAA
185221    AAAAACTCCC CCCAAATGAG GGTGAAATAA GACAATTTAA TACAGAGAAA AGAGGAAGGA
185281    ATTTATCTAG TCATATGTGA GAGTTTTATG ATACATTTTG TACTGTATAT GTGGATGTTT
185341    TCTATTTCAT TTAAAAAATC AACCGTGCAA TTAAATGGTA GATTGTCTTG CTTCTTTTTG
185401    ATTGACACAG TCATTAACTA AAATATTGTA GTATTTTTT ATCTCCCTGC CTAAAGGCAA
185461    TAAACATCTA ATCAGCAGAC TAGAACAATA AAAAATATTT TTTAAAGTC CTTTAGGCAG
185521    AATGATAAAA GTCCCTTAGG CATATTGAAA TTCCTATTTA TACAAAGGAA TAAACAGTAC
185581    TAGAAATTGT AACTATGTGA GTAAACAGAT AATATTTTTT CTCCATAAAA TGTGGTTGAC
185641    TATTTTCACA AAAATAGTTA ACAATGTAAT GTGTGATTTA TAGCATTTAA AAGTAAAACA
185701    GGCCGGGCAC AAAGGTTCGT GCCTGTAATC CCAGCACTTT TGGAGGCCGA GGCGTGCAGA
185761    TCACTTGAGG ACAGGAGTTC AAGACCAGCC TGGCTAACAT GGCAAAACCC CATCTCTACT
185821    AAAAATACAA AAATTAACCA GGCGTGGTGG TGCACGCCTG TAATCCCAGC TACTCTGGAG
185881    GCTGAGGCAC AAGAATCACT TGAATCCAGG AGGTGGAAGT TGCAGTGAGG CAAAATTATA
185941    CCACTGTGCT CCAGCCTAGG CAACAGAGCT AGACTCTGTC ACACACACAC ACACACACAA
186001    AAGAAAAGTG TATGACAACA ACAGTGCAAA AGAAGTGGAA ATGAAAATAA TGTTATTTTA
186061    TATAAGTGGT ATACTTTTAG ATGAACTACG ATAAATTAAT GATGTATACT ATAAACTCTA
186121    AGGCAACCAC TGAAATAATG AAACGAAGAA TTATGGCTAA CAAGCCACAA AAAGAAATAA
186181    AATAGAATGA GAAAAAATAT TTAAGTTGTT CAACAGATGG GAAAAAAAAG AGGAAAAAGA
186241    GAACAAAGAA CAGATGGGAC AAATGGGAAA GTAATAGCAA GATGATAGAC TTAACTCTAC
186301    CCATATAGAT TATCACACTT AAGGTAAATG ATCTAAATAC TCTAATACAA AAGCAGAGGT
186361    TGTCAGATTG AATTAAAAAA ACAGACAACA ACAAAAAAAA GCAAAAAAAG AGCCACAACA
186421    TGCTGCCTAC AAAAAATTCA CTTTAATATA AAGACACAAA TAGTCTAGAA CACCATCACT
186481    TTTAACCTTA TTTACTCAAA CCTCCTGATC CCTATTTATT TATTTATTTA TTTATTTATT
186541    TATTTATTTA TTTATTTATT TTTGAGACAG AGTCTGACTC TGTTGCCCAG GCTGGAGTGC
186601    AGTGGCACCA TCTAGGCTCA CTGCAGCCTC TACCTCTCGG GTTCAAGCGA TTCTCCTGCC
186661    TCAGGCCTCC CAAGTAGCTG GGACTATAGG CACATGCCAC CATGCCCAGC TAATTATTAT
186721    ATTTTTAGTA GAGACGGGGT TTTGCCATGT TGGCCAGGTT GGTCTCAAAC GCCTGACCTC
186781    AGCCTCCCAA AGTGCTGGGA TTACAGGCGT GAGCCACAGC ACCCAGCTCC TCTTCATTTA
186841    TTCTTGCTAC GCTTCCTCCA ATCCATTTTG TGCATTTGAT GATTTTGCCA GTAACTTCTT
186901    TATTTTTCTG GTAAAATTAC TTATGGGTCA CTGAGGACTG GGATGTTCTT TCTTCTAGAG
186961    GGGGTTTGTG TCTGCTTTTG CCAGGAAGCT GGGGTACCAC CAGTCAAGTA TTACTTTAAA
187021    CTCAATTCAT GAATTGAGAC TTTTTTTTTT TTTTTTTTT TTACGCAGAG TCCTACTCTG
187081    TCACCCAGGC TGGAGTGCAG CGGTGTGAAC ATGGCTCACT GCAGCCTCAA CCTACTGAGC
187141    TCAAGCAATC CTTCTGCCTC ACCATTCTGT ATAGCTAGGA CTACAGGTGT GTGCCACCAT
187201    GCCTGACTAA TTTTTTAAAT ATTTTTTTTA GAGATGGGGC TCACTTTGTT GCCCAGGCCA
187261    GTCTCGAGCT CCTGGGCTCA AGTGATCCTC CCACCTTGGT CTCCCAAAGT GCTGGGGTTA
187321    CAGGCATGAG CCTCTGTGGC TAGCCAAGAC TTTTTATTTT TTAGCCTAAA TGTGTATAAA
187381    AGTTGGCTTG TGGTTACAAC TTATCAGGAT TGATGATCTC TCTCTCTCTC TCTCTCTCTC
187441    TCTGTCTCTC CCCACCTCTC TCACATCCCT TGCTCTGCTG AGAAGCAGAG CAAACATTCT
187501    AGCAGTTTCC AGAGAGTAGG ATGGGATTAC TTCTAGTTTA CTTTTATCAT CCTTTGGGAT
187561    CGCAGTATTA CTGGGAGAAC ACAAGTATCT CTTATTAGAC ATACCACCTT TGTAGAATCT
187621    GGACTTTCAT TTTAGACTTT ATTTGTTTTC TACTATAAGC AATTTAAGTT ACAGATCTCT
187681    CTACACACTG TTTAAGTTGC ATCCCATGAA TTTTGATGTG CTTTATTGTC ATTATTATAT
187741    AGTACAATGT ATTTTGTAAT TTTTTGTGAT TTGTTTGGAG AGATTGATTA ATTAGAATGA
187801    TGTTTAATTT CCAAATATGT GTGTTTTTT CTACATTTCT TATTTTTATT GATTTCAAAT
187861    TTATTTCTAC TGTAGTCAGA TTTAATAATT CATTTATTTT TATTATTTTC ATTTTTTTAG
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187921    AGACAGGGCC TTTCTGTGTT GCCCAGGTTT GTCCCAAACT CCTAGTCCCA AGCAGTTCTC
187981    CTGCCTCAGC CACCCAAAGT GCTGGGATTA TAGGCACGAG CCACCCGTGC ACAACCAACA
188041    ATTCATTTAA AAAGTGGGCA AGTGAACTGA ACAGACATTT CTCAAAAGAA GGCATACAAT
188101    TGGCCAACAA ATATATGAAA GAATGCTCAA CATCACTGTA TTAGTCTGTT TTCATGCTGC
188161    TAATAAAGAC TTAACCTGAG ACTGGGGAAT TTACAAGAGA AAGAGGTTTA ATGGACTTAC
188221    AGTTCCACAT GGCTGGAGAG ATCTCACAAT CATGGTGGAA GGCAAGGAGG AGCAAGTCAC
188281    ATCTTACATG GATGGCAGCA GGCAAAGAGA GAGCTTGTGC AGGGAAACTC CCGTTTTTAA
188341    AACCATCAGA TCTCGTGAGA CTCATTCACT ATCATAAGAA CAGCATAGGA AAGACCCGGC
188401    CCATAATTCA GTCACCTCCC ACTGGGTTCC TCCCAGGACA CATGGGAATT GTGGGAGTTA
188461    CAATTCAAGA TGAGATTTGG GTAGGGACAC AGCCAAACCA TATAAATAAC TAATCATCAG
188521    GGAAATGCAA ATCAAAACCA CAATAAGGTA TCATCTCACC CCAGTTAGAA TGGCTATTGT
188581    CAAAAAAACA AAAAATAACA AATGCTGGTG AGGATGTACA GAAGAGGGGA CTCTTATGTC
188641    CCACTGGTGG AAATGTCAAT TAGCATAGCC ATTATGCAAA ATAGTATGGA AGTGAGGTAG
188701    GTTACATAGG GTGGTCACAG CCTCCCTTGA AAGGAAACAA GAAACTTGTC AAATTGATGG
188761    AGAGAACAAA TCTCTTGACA TTACACAAAC TGCATCTGGG GCTAGTGGTT AGAATATCCT
188821    CAGTCAAGGA GGTAGAAGAG CAGGAGGGAA AATCCCTAAG TTCGTGCAAG TGCAGAAACC
188881    CACAAGCTGT GTTCTCAGGT TGACATATAC TCATTTTAAT AGTAAGAAAC ACACCCTTGG
188941    GTAGAGAATT AAAATGCTAA TAATACATGT GATGTATGTA CTAGCGTGTA TGGCAATATT
189001    GCATGCACAT TCAAGAGACC ACCCAAAACA TATTTAACAA CAATGCCCAT TCCCACCCCC
189061    TCATGGATAA TCACGTAGGA CTCCCATAAC GGGAGTTTCT TCAGTGTCAA TTGGTGCTGA
189121    AGTAGCCGAC CCTGACTCTG CTATCAGCGT GTACTTTCAC CTTGCAATAA ACTCCTTTGC
189181    CTACTTTTAC TTTGGACTGG CTTTCAAATT CTTTTGTGCA GGGAATTCAA GAATCTGAAC
189241    CAGCCTACTG ACAACAGAGG TTTCTCAGAA ACCTAAAAAT AGATCTACCA GATGAGGCTG
189301    AAAATCTGCT ACTGGCTATT TATCCAAAGG GAAGGAAATC AGTATACAAA GAGACACCTA
189361    CATCCCCATG TTTATTGCGT CACTCTTCAC AAGAGCTGAT ATATAGAGTC AACCCTAAAT
189421    GTTCATTAAC AGACAAATGG ATAGAAAATG TGGCATATAT ACACAATGAA ATACTATTTG
189481    GCCATGAGAA GAATGCAATC TTGTCATTTG TGGCAACGTA GATGAAACTG GAGAACATTA
189541    TGTTAAGTAA GATAAGCTAG GATTGGAAAG ATAAATACTA CATGTTATCA CTCATATGTG
189601    AAAGTAGAGA AAAATTTTA GCTCATGGAT TTAGAGAACA GAACTGTGGG TACCGGAAGC
189661    TGGGAAGGGT AGCAAGGAGG GGAGGATAGG GAGAGGTTGG TTAATGGTGA CAAAATTACA
189721    GCTAGATTGT AGAAATGAGT TCCGGTGTTC TGCACCATTG TAGGGTGCAT ATGGTTAACT
189781    CTCATTTATT GTATATTTTC AAAAAGCTAG AAAAGAATTT TGAATACTCA CAACAAAATA
189841    AATGATAAAT GTTTAAGGTG ATGGATATAC TAATTACTCT GATTTGATTA TTACACATTG
189901    TGTACACATA TAAAAATATC ACTCTTTATC CCGTATATAT GTACAGTTAT TATATGTCAA
189961    CTAAAAATAA AAGAAAAAAA GAATATGATC TATCATGATG TATATATCAT GTGTACTTGA
190021    GCAAAATGTG CATGCAGATA TTGTGTATAA TGTTCTATAA ATCAATTAGC TCAAGATAAT
190081    AGATAGGATT GTTCAGATCT TCTGTGTCTT TACTGATATT TTGTCTAGTT ATTGCATCAT
190141    TACCAAAAAA AGGGTGTTAA ACTCTCCAAA TGTGATTGTA GAATTGTCTA TTTTGTCTTT
190201    TCTTTTCCAT TTTTACTTTA TGTATTTTGA AACTCTGTTA TGACATTTTG CTATGTATTT
190261    TAAAACTTCG TTATGTATTT TGAAACTCTG TTGTTAGAAT CATACATTTA TGATTATTAT
190321    GTTTTCTTGA TGAAATGACA CTTTTCTATT GTCATTGTTT TTGTTTTTTC TGAAATGGAG
190381    TCTCACTCTG TTGCCCAGGC TGGAGTACAG TGGCACAATC TTGGTTCACT GCAACCTCCA
190441    CCTCCTGGGT TCAAGCGAGT CTCCTGACTC AGCCTCCAAG TAGCTGGGAT TACAGGCATG
190501    TGCCAGCATG CCAAACTAAT TTTGTATTTT TATTAGAGAC AGAGTTTCAC CACGTTGGCC
190561    AGGCTGGTCT CGAACCTCTG ACCTCAGGTG ATCCGCCCAC CTCGGCATTT TTATTTTATT
190621    TTATTTTTTT GAGACAGAGT CTCACTCTGT CACCCAGGGT AGAATGCGGT GGTGTGATCT
190681    TGGCTCACTG CAACCTCCGC CTCCTGGGTT CAAGCAATTC CCATGCCTCA GCCTCCCGAG
190741    TAGCTGGGAT TACAGGCACA TACCACCATG ACTGGCTAAT TTTTGTATTT TTAGTAGAGA
190801    TGGGGTTTTT CTATGTTGGC CAGGCTGGCA ACTGACTCCT TTAACAATAC AAAATATCAC
190861    TCTGTCTCTG GTAACACTCT CTGTCTTAAA CTCTATTTTA GCTGTTATTA TTATAGCCAT
190921    TTTAGTCTTT TTATGCTTTC TGTTTGCATA GTGTATATAT TTTAATATGT TTATTCTCAA
190981    GTTATCTGTG TTTTTATATT TAAGATGTTT CTCTTCTAGC CAACGTGTTT GGTTCTTGCA
191041    TTTTTAAGTC GATTCTAACA ATCTTTGCCT TTCAATTGAA ATATTTACAC CATTAACATC
191101    TAACATTAAC ATTTATTTTT CTTTCCACAG TACACTGGCT AGCATCTCCC ATATAATATT
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191161   GAACATAAAG  TGTGATAACT  GACATCCTTA  TTTCATTCCT  ACTCTGAGTG  GAAAGGGCAG
191221   GGGTGGAGAA  AGCATTCAAC  AATTTGCCAT  AATTATAATG  CTTTTTGTTA  CACTGTTTTC
191281   TTCTGCATTA  AAAAATATCA  TTACATTTTG  CATGAATTAT  TAGGAGAAAA  TATTTTCCAA
191341   TTTTCCTGGA  AAATGCCATA  ACCACGTCTC  TCAATTTTGT  TTCCATCTTT  CTTCCACATT
191401   TTACATAACC  TACATAAGAG  ACACATTATC  AAGTATATTT  TACATGGCTT  CTCAGTGTCT
191461   TCTCTGTCTG  CTAACAGGTT  TACCAAGAGA  TGGCACTCTT  GTATTCTGG   TGGCTATGTC
191521   CATATCGTTT  TGCCTTTAAG  ACAGCGTAAC  TACTTCTTTC  ACCAGTATTA  AAGACATGTA
191581   CATTTGATCT  GGTTCTTGTG  GATGATTTTA  AATGACTCAA  GCTAATAATC  CTAATTTTAC
191641   CTAAACACTC  CATTATTTTA  AAATGTATTC  CTTTATGCCC  ACAATAAACA  TTTATTGACA
191701   TTAGGCTGGA  CATTAGGCTT  CTCTATGGCA  GACATTAGGC  TGGACCCTAG  CCATATATCT
191761   ATTGAGGGAA  AAAAAATTAT  TTTCTATATA  AGTTCCAGA   AAGCCAAGAT  GTGTTTTAAA
191821   AACAAACAA   AACATTACAT  TCTAAATGCT  GTAACAAGAT  AAGAAAAAGT  GTTGAGGCTG
191881   AGAGAAGAAC  AAAGCAGCAA  GCAACTCCTG  GAAGGACCAC  TGCTGCAGAG  GTAATAACTG
191941   GTGAACCATG  TTTTGGAGAA  GGAAAAGGTC  ACCAAGAGAA  GGAGGGGGTC  CAGGGTGTTC
192001   AGAAAGATTG  CATGCATAAA  GATCAAGGGT  AATAAAAAAA  ATTCCGTATT  ATGTAAATGT
192061   GAAGTTCCAG  GACCATGAGC  TTGGAGAGCA  TGAAGTACAG  GAGGAGGGTT  GGTTTCAAAT
192121   AAATCTGGGA  ATGAAACAGT  GAAGCCTCTG  GCAGAACTCA  CATCTCTTTC  CTCCCCTCTT
192181   CCTTGCACAT  TCCCTTTATG  GAGTAATTGC  AGGGATGGGA  AAAGTTCAAA  ACCACCACTG
192241   AGCCTAGGAA  GTGCTAGGGT  AAAGTGGAGA  ATGAACCTGC  GTGATTTGCT  CATCCTAAAC
192301   TAGGTTCTTC  TAGGAGAGCC  CTTCCCCATA  AAATCTGCCC  TCCTCGAAGG  GGCCCAGACA
192361   GCCTAAGCTC  ACCTCCCAAA  GACCCCTTAC  TTGCTGACTG  AATCTGATTC  CACCCAGACA
192421   TGGCCTAAAA  CCCTTCCATA  ACTCTATAGC  CAAATTCAAT  TTTAGACAGG  CCTCATACCA
192481   ACCTTTCTTC  CTCTAAGTCT  GCCACCCTAG  GCAATTCTCA  ACATTCTCTA  CACACTTTGG
192541   GGCCATAGAC  GTGCTACCAA  GTCTCCAGAC  CTAGACCTGA  TGGAGCAGTG  CTGTAATGAG
192601   ACGACCACTG  GCCTTTGAAC  CAGACCCTTC  TCTGTGGCTC  CTATGCATCT  CCAACCTGTT
192661   TTGAGCACTG  CTGCCAAGAC  ATCTTTGGCA  CTTTGTTGTG  AAGTTTTAAA  ACTGAACTAA
192721   TCTACAAAAC  ACCTAACCTT  TAAAAATTCA  TTGTCATTTC  ATATCATGAA  AGATAAAGAA
192781   AGGCCAGGAA  ACTGTTCCAG  GTTAATAGAG  ACTAAAGAGA  TAGCAACCAA  ATGCAATTTG
192841   TGATCCTGGA  TTGAGGGGAA  AAAGTGTTGT  CAGAGACATG  ATTGGGACAG  CTGGTAAAAT
192901   TTGAATTTGA  ATTTAAAGAT  AAAGTATTGA  GTAATATAGG  AAGATGATTA  TCTGCAACTT
192961   TCAAATGTTT  CAGTAAGTAT  ATATATATAT  AAAGAGATAT  AAAGACATAT  AAATAAATGG
193021   ATAGGTAGAG  AAAAAGCAAA  TGTATAATAT  TAACAATCTA  GGTAAAAAGT  ATATGAGTGT
193081   TCTTTGTACT  GTTTTTCTGA  TTTTTCTATA  TGTTTGAAAT  CATTTTAAAA  TAAGAAGGTT
193141   TTTGGGTTTT  TTTTGTTTGT  TTTTTGTTTT  TAGAGACAGC  ATCTTATTCT  GTCACCAGGC
193201   TGTAGCTCAG  TGGCCCAATC  ATTGCTCACT  GCAGCCTCAA  CTTCCTGGGC  TCCAGTAATT
193261   CCCCCTACCT  CAGGCTCATG  AGTAGCTGGT  ACTTCAGGTG  TGCACCACTG  CACTCAGCTA
193321   ATTTTTATTT  TTTAAATTTT  TGTAGAGATG  GCATGTTGCT  ATGTCACCCA  GGCTAGTCTC
193381   AAACTCCTGC  CCCCAAGTGA  TCCTCCCACT  TTGGCCTCCC  AAAGTGCTAG  AATTATAGGC
193441   ATGAGCCACT  GCACCCAGCC  CCAAATAAAA  AAGTATTTTA  TTTTAATTAA  CTAATTAACT
193501   TTGAGTCAGA  GTTTCACCCT  TGTCACCCAG  GCTGGAGTGC  AATGGCATGA  TGTTGGCTCA
193561   CTGCAAACTC  TGCCTCCTGT  GTTAAGCGA   TTCTCTTGCC  TCAGACTCCT  GAGTAGCTGA
193621   GATTACAGGT  GCCTGCCACC  ATGCCCAGCT  AATTTTTATA  TTTTTAGTAG  AGACGGGGTT
193681   TCAGCATGTT  GGTCAAGCTT  GTCTCAAACT  CCTGACCTCA  GGTGATCCAC  CCACCTCCGC
193741   CTCCGAAAGT  GTTGATGAGC  CACCACACCC  GGTCTAAAAA  GTATTTAAA   ACCACAGTCC
193801   CACTCTACCT  TGTCCTACAC  TACCAGGGGC  TAGGATCACC  CCATGTCTTC  TAGGCTATGA
193861   GATAGAGGAA  TCCAAGGAAG  AAGATAAGCT  ACTTGGTTCC  TCTATAGGGT  CTTGTGTGTG
193921   CTCTCATGTG  CTCTCTCTCT  CTCTCTCTCT  CTCACACACA  CACACACACA  CACACACACA
193981   CACATGAATA  CCAGAGCTAT  CACTTTCCCA  GTCTAGTACT  CATCTCATCC  CAAGGGTTTT
194041   GTGTTGTAGT  GGTTTGCTCA  TTTCTTTGTT  TTGTTTGTTT  GCTTGGATTA  TTCTTTTTCT
194101   CTTTTTGCAG  CTGAAGGGAG  AATTTCCAGG  CCAGCCCTTT  GGCCATTAGA  GTTACAGTGC
194161   CTCTATTCAG  GCTTCATAGA  GAGACCTGGG  ATTCAGTAGT  GGGGGGCTTT  TATCCAGTTC
194221   AAAATAATGC  ATTCTCACCA  AGATGTACTT  TGAAATAAAA  CAATACTAAA  ACACAAAATT
194281   TTATTTATGC  TGAACATTGA  ATCACTTTTT  TCTGTATTTT  GTGTAGAAAG  TTATACACAC
194341   ACAAACACAT  TTGCTCCTGC  TTTGTTTATT  GGCCCAGGGG  TATGTTTGGT  AATACTTCAT
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194401   CAGGCATGAG TAGTACGTCT TGGAAGGTGT GGTCTAAAGC CTAGACTCCT ATCTGCTTCC
194461   TTCAGCATTC TCCAGTGTAT CTGTCATCTG TCTACCTTAG GATAGGGGTC TCCAGAACTT
194521   CCATTCACAT TTAGAAGAGG GCAGCGGCTT TCTATGGAAA ATATGAACTC TCATTCATCT
194581   CTATTCCTTC TTCTAGCTAT GGTCCAGCTC AGCTGTTTGG AATAAAGTAT CTATATGAAG
194641   TCTGCGAATG GTTCTCAGAC TGGTTGAACA TTAGAATCAC CTGAGTACCT TCTAAAATTC
194701   TTATTACCCA GGGCATATCT CAGAATGAGT ACCGCAGGGT AGGGATAGGA TTAGGGATCA
194761   TGATCTCTGG AGTCTGGTTT AGGCACTAGT GCTGTTTAAA ACTACGTTCA TGAGGTGGAG
194821   GTTGCAGTGA GCCGAGATGG CGCCACTGCA CTCCAACCTG GGCGACAGAG TGAGAGTCTG
194881   TCTCAACAAA ACAAAACAAA AAAAACCAAC TACCCTTGTG ATTTGAATGT CCATCCAAAA
194941   TTGAGAACCA TTAGGTAAGG CCAAGCTGTA TAATTAAAGA GCAGTTTTCA TTTGTCTGGT
195001   GTGGTGGCAG CTTTTTGATA AGGGAAGTAT TGTTGCCATC CACATACCTG AGCCTCACTC
195061   CTGAGAACAC TGGTGTGTAT GTTGCTAAAA TTCCCCAGGT GATTCTGAGG TTCCTTCCTG
195121   GATAAAAACC ACTGACCCTG GGAATGTACC CACTGCCAAT CTCCTGCGTA AACCTTGGAT
195181   ACTGGGAAGC CTACAGTTGA AAATATTGGG CTTGAGATCC TGAAACAAAT CTTGTATTTC
195241   ATTAAGACTA ATATTTGGTA CAGTGCAGCA AATCAAGGGA ATTTTGGTGG CTGAGTTCTT
195301   TTAGAACTTT TGCATTGAAA TAGGTTCAAG CAGCAATAAG TTAAAACTAC AACCTCAGCT
195361   AAAGGATTAA AAGACACGTG AGCTGGGTAG GATGAGGTCT AAGGTTGGGT GTGGCGGCTC
195421   ATACCTGTAA TCCCAGCACT TTGGGAGACT GAGGTGGGTG GATCACTTGA GGTCAGGAGT
195481   TCAAAACCAG CCTGGCCAAC ATGGTGAAAA CCCATCTCTA CTAAGAATAC AAAAAAATTA
195541   GCTGGGCGAG GTGCCAGGCA CCTGTAATCC CAGCTACTGG GGAGGCTGAG GGAGGACAAT
195601   CACTTGAACT CAGGAGGCAG AGGTTGTAGT GAGCTGAGAT CGCACCACTG CACTCCAGCC
195661   TGGGTGACAG AGCAAGACTC CATTTAAAAA AAAAATAATA ATAATAACAA TAATAATAAT
195721   TCAGACATAT CCAGGCATCA AACAGATACC TGGGGCAGAT GAATAGTCTT GAGATTCAAG
195781   TCACACATGA AATTTAGGTG GAAAATGACA TTGGAGAAAT TTGAGATTAT GATGAATGGA
195841   AATTTTTCAA AGAGGAATTT CAGGCTCTGT TCTTGAGGGG ATAGATGGAC TTCCAACAGC
195901   AATAACACAG GATTAATGAG GACTTGGGAT GTTACATAAA TTAGAGATGT TAGATGGATA
195961   AAGAGATAAA AGTACTCTCT CTAAGAACAT GGGACCAGAG ATAGGCTCAC TTCTAACCAT
196021   CAGATATAAC TAGCAGACTA AACGGTCTAA AAATAAAAAT CATGCCCCAC TCCTGCTTAA
196081   GACATTTTAA TTACTCTCAG TAACTCTTCA GTTTTTCTAC TGTGTTATCT TTAACTACAG
196141   GGTTGGTCTG GGTGTGCAAC ACAAGAAAGC CTGGCATATA CATGGATTCA AGTGTATGCC
196201   ATGTGCAGGT ATTCTTTCAT GTACTATTTC ATGTATTCTT TTTCACATCT GTTTTTTCCT
196261   TCATTGAAGT CAATGGCTGA TATTAGATTC TACTATTCAT GTGTACTAGT TATATATAAT
196321   TGTTACAAAA CAAATTAGCA AAAACTTAGT GGCTTAAAGC AACACACATT TATTATTACC
196381   TAAGGTCTGT GGATAGAAGT TCTGACATGG CTTAACTGGG TTCCCTGCTT CAAGCCTCAT
196441   GTGGCTGCAA TCCAGGTGTT GGCTGAGTCT GAATTCTCAT CAGAGGCTTG ATTGTGGAAA
196501   TTTCCACTTC CAAGCTCCCT CAGGTTTGTT GAAAAATTCA GTTCTTTGCA CCGGTAGAAG
196561   CTTCTTGGTA GAGGCTGATT CAACTTCTAG AGGCTGTCTG CAGTTCCTGT CACCCAGGGT
196621   GGAGTGCAGT GGAGCAATCA TAGCTCACTG CAGCCTTGAC CTCCCAGAAT CAATCTGTTC
196681   TCCCACCTCA GCATCCTGAG TAGCTGGGAC CACAAGTGTG TGCCATCACA CCTGCCTAAA
196741   AAACAAACAA ACGAAAAAAA ACCCCAGAG AACTTTGTAG AGACAAGCTG GTCTGGAACT
196801   CCTGCGCTCA AGCAATTCTC CTGCCTTAGC CTAAAAGTTC TGGGATTATA GGTATAAGCC
196861   ACCATACCTG GCATATGGCA AGTCTTGAGC AGGACAAATA CAGATGATTT ATGTCTGTCT
196921   TCCATGGTAT TCTAGGTTAT TGTTGAGATG GTCCTCTATT GTCTTGTTCC ATCTATTGAT
196981   TAGATAAAAC GTTGTTCCTT CTGTTATTTT TCAACAGTAG CTTTTATGTG TCTCTCTTTA
197041   TCTTAAAATT CTAACCAAAG AGCTGCTCTT TTCTTGGTGT ACTTTACCTT TGGTTGATCC
197101   TTCTTAACCT CTTCTTGCCC TCTGGGGCCT AAGATGAGGG CTGTTATCAG ATGTGAGTCT
197161   ATGGGAAAGC AAGCAAGAGG TTCTTCAGCC TCCGTTCAGC CTTAAATGTC TAGGTAGAAA
197221   TCAGTCATGG CCCTTCCAAT GTGGTACAGA CCAGATCACA GAGACAGGGG TCTCAGCCAA
197281   GGTCTTGTGG CCTAAGCCTT ATAGAAATAA TGAGTGTTTA CTTACTTGGA GAACTCCCTT
197341   GGAATATCTT TTTTTGTGAA CCTGAGGCAA CTTTTGGTGA TTTCTTGATG TCTTGGGAAT
197401   CTTGGTCTAG AGCCATTTCA ACCCGATTTC TTTTCATGTC AGTGGCATTT TGTGACCAGA
197461   TAGTAAATAA GTTCTATGAT GTTCACTCAG AGAAATACAA TGACTTATGA TGCGAAGCTT
197521   CTGTGGTTCA GCCCTTACTT CATCTTCATT CCCTCTTATC TGCATCTGTC TCCTGCTTGG
197581   GAACAAAGT CTGGCTTCAT TCTATGACCC CCACGTTGAG TTTCTTAGTA GCACTTACTT
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197641   TTCAATTAGG AGTGTCCTCA CTTCTATCCG TCAGACATAA CTAGCCGACT AAACAGTCTA
197701   AATATAAAAA TCATGTCCTA CTCCTGCTGA AAACATTTTA ATTACTCCCC ATCATTTAAT
197761   TTTTTCTACT GGGTTATCTT TAACTTCAGA GTTGGTCTTG TGTGCAACAC AAGAAAACCT
197821   GGCATATACA TGGATTCAAG TGTATGCCAC GTGCATGTAT TCCTTCATGT ACTATTTCAT
197881   GTATTCTTTT TCACATCTGT TTTTTCCTCT AAAATTTATT TCCTTTTAAA AATGAAAATT
197941   TTGCATTTGA CTAAATTTGT CAAATTTAGT CAAATTTGTT TAAAACCATT TTTAAAATGT
198001   TTCCCGAAGT TTTGAGTGAA GTTAGTACTT CAGAAAAACT GTTTTGTATT TTTCCTGTGA
198061   CCTCAGTGCA CTGCTGTGCA TTTCCATTTC TGCGTCCACA CACATTTGTT TTGAGGAAAT
198121   ATAGGAACGA CAAGATAAAG TTCAAGCTCC TGGACATTGC ATAAAAGACC GTCATGACCT
198181   GGTCCTGTTG ACTTCCCTAG ATTTCCCGCT ATTTCCTAAG TTGAGATTTT TGGTTTGGAT
198241   GCTTTGTGTT TTCCTAAAAT CAAAATAGGT TTTTGCCTTT TATGATTATA CAGTAAATAA
198301   ATGCTATTTG TGTGAAACTT TAAACAATAC AAAAAAAACC TAAGGAAGAA AGTCAGATTC
198361   ATCTAAAAAT CCTTGTGGCC AGAATTAACT ACCTTAGTTA CTATTTTCTC TATCTCTCTC
198421   TCTCAATGTA TATTTGGTGT AGGTATAGGG GTGTGTGTAG TGTGTGTGTA TGTATATATC
198481   TGTTTCTATT CCTGTATGTG GATGTGCACA ACGCATCCTG CTTTGTACAC TACAGTACTA
198541   GCATTTTTCT AATGTAATTC AATATTGTTG AAAACATTTT AAAAAAGCTT GTATATATAC
198601   ACACACATAC ACATACATGC ATGTATGTAC ATATACACAT ACAGACAAAA ATGTATCCTA
198661   TGTATATTCA CACATGTATA CACACTCACA CATACATAGA GTTTTACATC CATAGTTTAT
198721   AAATGTTGCT TTTTTTTGGT CACCTTTTTG CTAAGTCTTA CACTTTTTTT TTTTTTTTTT
198781   GAGACGGAGT TTTGTTGTCA TTGCCCAGGC TTAGTGCAGT AGCGCGATCT CACCTCACTG
198841   CAACCTCGAC CTCCCGGGTT CAAGCGGTTC TCCTGCCTTA GCCTCCTGAG TAGCTGGTAC
198901   TACAGGTGTG CGCCACCATG CCTGGCTAAT TTTTGTAGTT TTTTTATAGA GACGAGGTTT
198961   CACCATGTTG GCCAAGCTGG TCTGGAACTC CTGACCTCAA GTGATCTGCC TGCCTCAGAT
199021   TCCCAAAGTT CTGGGATTAC AGATGTGAGC CACTGCACCC GGCCAAGTCT TACACATCTT
199081   TTTTTTACCA CTAAACTGTT TACCCAAACC TGATAACCCA AGTCAACAGC TATTATGGCT
199141   CACACAATCT TATGTAAACA AAGATACAGA TATATAGAAT TTTCTTGATT AATATTCAGA
199201   AAAAAATGGA GTCCCTTTAT ACGTCCTTAG TATCTGCTTT ACTCATTTAA AAATGTATTA
199261   CATTATATGA AAGTATTCAG GTCAAATGTT ATAGATGTGA TTCATTCTTT TTAACTGTGT
199321   TATTTTTCTG CAATGACTAT GTATCACAAA GTACTCAGTC TTCCACTGAT GAAAATTTGG
199381   GCTATTTCCA GTTTGTCTTC CATTTTTCTT TCTTCCTCTT GGATTTTCAC TCAATGTGTT
199441   TACTAATTTA GGAAGAATCA ATAGTTTTTA TGGTATTACT TCTCCCATTC AAGAATATAG
199501   CATATGGTAT AGTATAGTAG AGTACTTAGT TTAATTTAGC CAGATCCTGT TTTCTGCCCT
199561   TTAATAAAAT TCTATCATTT TCTGCCTTTG AGTCACATTT TCCTTGTTCA TATAATTCTT
199621   AAAAAATGTA TAGTTTTCAT TCTAAGGGAA CATAAAAACT TCTTTCCATT TCTATTCCTG
199681   TCTAGTTAAT TCTACTATTG GGAAAAGTAA CTGTTAAAAA AAATTCTTAT CTTTCCAGTC
199741   AGTTCACCAC ATTTCCTTTA TACCTTTGTA CTTTAATCCC CAGTCATGTT GAACACTTCT
199801   TATTCCTCAC ACCAAGCCTC AACGGGTTTG CTCTTTCTGG AAGGTGCTTC CCCTGTATTA
199861   CTGACTTATT CATACCACAC ATGGAGACTG GCGCAGCCCT GTTCTGCCTG GGAAGCCTTC
199921   CCCTGATACC CCCAGTTGGC AGGAGTCTTC ATTTGTTCTT TTCTAGTCAC CTGTGCAAGT
199981   TTGTATTGTT CATGTTTATC ATCCTTCATT CTAGTTGTCT GTCTCTGTGT GTGGTCTCAT
200041   TCAGTGGACT CTGAACTCTT ATGAAGTCAT GTCATGGGTC AGATCTTAAT AAATTAATAT
200101   TGTCGGAAGC TAATGTCATG TCTAGAATAC AGAAAATTTA TCAAAAAAAA ATATAGTATG
200161   TTGGCTGGGC GCAGTGGATC AAGCCCGTAA TCCCAGCACT TTGGGAGGCC GAGGCAGGAG
200221   GATCACATGA GGTCAGAAAT TCAAGACCAG CCTGGCCAAA ATGGTGAAAC CTCATCTCTA
200281   CTAAAAATAC AAAAAGTAGC CAGGCGTGGT GGTGCCCACC TGTAATCCCA GCTACTCAGG
200341   AGGCTGAAGC GGGAGGATCA CTTGAACCTG GGAGGCAGAG ATTGCAATGA GCTGAGATCA
200401   TGCCACTGCA CTCCAGCCTG GGCGACAGTG AGACTCCATC TCAAAATAAT AATAATAATA
200461   ATAATAATAA TAATAATAAT AATTGTATGG AATTGAACTG CTCTGATTGG AAATAGCTGT
200521   TTTTTAAAAA ATTATTATTT TTTAAGTTCC TGGGTACAAG TACAGGATGT GCAGGTTTGT
200581   TACATAGGTA AACGTGTGCC ATGGTGATTT GCTGCACCTA TCAACCCATC ACCTAGGTAT
200641   TAAGTACAGC ATGCATTAGC TCTTTTACCT AATGTTCTCC CACACCCCCA CCCCATCCTC
200701   CCCCAACAGG CCCCAGTGAG TGTTGTTCCC CTCCCTGTGT CCACATGTTC TCATTGTTCA
200761   GCTCCCACTC ATAAGTGAGA ACATGAGGTG TTTGGTTTTC TGTTCCTGCC TTAGCTGTTA
200821   ATGTCAGGCC AGAGAGGCTT AAATTTTTAA GGATCTCTGG ACTTTTCTTC TACATTACTC
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200881   TTGATGTTTA TAAATGTTAC AACTTCTTTA ATTTCATTTA ATGTATACCT TATTGAGTTG
200941   ATTTAACTGA GTTAACTTTG TTATATGAAA ATCATGATTG GGAGTGAGGG GGTTAAACCA
201001   GCTACAGAGA TCTTGATTGT TGGTGGTGAA GCAATGCAAG AATTCATTCA TTCAGTAAAC
201061   TAATGTTTAT TAAGCGTGTA CTGTCTTAGT CTGTTCAGAC TGCTGTAACA AAATATCATA
201121   AACTGGGTGA CTTATAAACA ACAAAAAATT TATTTCTTAC AGTTCTGGAG GTGGGAAGTC
201181   TAAGATTAAG GCCCTGGCAA ATTTAGTGTC TGGTGAGGAC AGGTAGCCAT CTTTTTGCTG
201241   AGTCCTAACA TGGCAGAAGG GTTGAATAAA CTTCCTTGGG TTTCTTTTAT AAGGACACTA
201301   ATCCTAGTGA TGAGGTTTCT GCCCTCATGG TATAACTACT GCCCAAAGAC CCCTCCTTCT
201361   AATATTATCA CTTTGTGGGT TAGGATTTCA ACATGAGTTT TGAGAGGATA CAGACATTTG
201421   GATCATAGCA CACACCATAG GACAGACACT GTGCCAAGAA TTGTGGATAT AGTGATTCTC
201481   AAAATGAACA AGATCCCCTC AGAGAGCTTG CAAAATCCAG CTATAAAATT ATGCTTTTTA
201541   AACAAATTAT GCAGTTTGAA AAATCTACTC TGAATCTTAC TTGTGGCATT GAATACTTTC
201601   GGCCACTCTT TCCTTATTAT ATTAAATATT TACTCTTGTT TGGGGGATCC AGTCTCACCT
201661   ACTTTTTCTA CCAGAACTGG TATCAGCTCA TGCTCTGCCT TATGCAAATT AAGAAAATAT
201721   CATACCTTTT GGGTAAATTA AGCCAAGAAA GTTCTCCTTT CTTCTCTTTC TCTCTTTCTT
201781   TCTTTCTCTC TTTCTCTTTC TTTCTTTCTC TCTCTCTCTT TCTTTCTTTC TTTCTTTCTT
201841   TCTTTCTTTC TTTCTTTCTT TTTCTTTCTG ACAGGGTCTT GCTCTATTGC CTAGGCTGGA
201901   GTGCAGTGGT GCAATCTCAG CTCACTGCAG CCTTGAACTC CAGGGCTCAA GCAATCCTCC
201961   TGAGTAGCTG GGACTATAGG CATGTGCCAC AACATCAAGC TAATTTTTGC ATTTTTTTGT
202021   GGAGACGGGA TCTCCCTATG TTGCTAAGGC TGGTCTTGGA TTCCTGGGCT TATGCGATTC
202081   TCCTGCCTCA GCCTCCCAAA GTCCTGGGAT TACAGGCATG AGCCACTGCC CCTGGCCATT
202141   ATAACTATTT TCATTGGCTT ATCAGGCACA TGATAACTAT AATAAATCAA TAACCAGAAT
202201   TTTTAAATAA AGAAAGGAAG GAATTGTTTC AACTCTTCCT GCTACCCCTC TATCCCTCAA
202261   AAGGGTAGGC TGAATGTTGT CCTCCAAAGA TATCCATGTC CTAATCCCCA GAACCTGTAA
202321   ATATATTACC TTATATGACA AAAGGGACTT TACATGTTTA ATAAGTTAAG AATTTTGAGA
202381   TGGGCAGATT TTCCTGAATT TTGCAGATGG GCCCTAGTGT AATCACAAGG GTCCTTATAA
202441   GAGACAGGCA GAAGAGTCAG AATAAGAGAA AAATACTTCA AGATGTTACA CTGCTGGCTT
202501   TAAGGTGGAG GAAAGGCCAA GAGCCAAAAA ATGCAGTGGT CACTACAAGC TGAAAAGAAA
202561   AAGAAATGGA TTTTCCCCTA AAGCCTCTGG AGGGGGCACA ACCTTGCCAA TACCTTGATT
202621   TTGGCTCAGT GAAACCCATT TTGGACTTCT GACCTTTAGA ATTGTAAATA AATAAATAAT
202681   TTTGTGTTGT TTCAAGCCAT CACAGTTGTG GTAATTTACT ACAACAGCAA TAAAATAGAA
202741   TTAAATACAG AGATCTGAGG AGTTGAGTAG GATAAGCCTA CTCCAGCAGG TTATTTCGGG
202801   AGTATGGTGA GACTCACTAG GATGGCGGAA CTCAATTAAG GAAGTCTGAA GCTGATAAGC
202861   CAGAGAGGGA AGGCTCTCAT TTCATTTTAT AAGGGTTGCG TCACACTAGG AAGATCCAAT
202921   AGCAACCACA GTCTCAAAAT TAATGATTAC AAATAGGACA CAATTCCAAG AGTCGGGAGC
202981   CAAGCAGAAA ATGGATTAGG GAAGACATGG ATGATATGAA ACAGGAAGGA GGGGTACAAG
203041   GCAGCTTCCT GGGAAGTTGC CAGGGCAGTC ACAGTTCACA TTCATTAGGC TGTGGGCACC
203101   AAATGCATAT GGAAAATCTA GCTGACTTAA CTGAACTCCT GAAGAGGAAT GAACACCTCA
203161   TTTATTGAGG AGCTACTACC AATTAGAATA TGTATTTCAT TTGTTCAATA ACCCCATGAG
203221   TACAGTAACA CAATCCTTGC TTTACTAAAG CGGAAGCCAA TTCAAAGAGG TTCAGTGACT
203281   TGTCCAAGCT CAGGGAAAAC ACTAGGAAGT GAATATGGGT CTGACTCCAT CACTGATTTC
203341   AGGAGCCCTG CCCTTTCCTC CACACCATGC CCCCTTGCTT TCAGAAAAAA AGGCTTGTTG
203401   ACTGAATGGT TGTATGCACA GTTCAAAGCA GAAACACACG ATGACATCTT TTGAGATACT
203461   CTAACAGTGA GAACTTGAAA ATGAAGTTAA AAATTAAGCG GCAAAACCAA GCCGAGGCTT
203521   TCTGAGAAAG TGGGGCCAAA CCTGTTGCCG TCTGACTGCC ACGTGGCTCA CTATTTATCC
203581   CTGTAAAAAT CTGCAAAAGT ATTTGAAAGG GAAGAAGGGA CAGAAAACTC CCTCCTTTTC
203641   CAAGTTAGCC TTATAGTCTA GGGCTTAAAA TACTGGTTTA ATGGTGAAGG TAAGTGCTTT
203701   TCTTCTTTTT GGGTAGAAGG ATTATTACTA ACTTACCAAA GGTCCATTAA GGGGAGGGAA
203761   CAGTTTTAGG AGAAGTCAGA GAAAGACAT TAACAGCAAC ATAAGGATCT CCATCTGGTA
203821   ATATTGCCTA ATTCCAAAAT GAAGAGACTC TCTGAAAAAG ATAACTGATT CAATGAAGAC
203881   CCTAGGGCAA GGCTTGAGAA GCCACTGGTA CCAATGGACA CTGTGGACAA TGGTCATTTC
203941   TCCAAGGACG CTGTGAGTAT TAACTGTGAT GCTGTGATTA GTCAGACTGG GATTGGCTGT
204001   GGAATGAAAT ACTGATCAGA ACTGACAAGA TTTGTGTTTG GGACTGTGGC TAACGAGTCT
204061   TTTCAGACTT CTATATGAAT TTGAAATGGT CTCTCAGGAA AAGGAGAACA TGGCCGGGCC
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204121   TGGTGGCTCA  CGCCTGTAAT  CCCAGCACTT  TGGCAGGCTG  AGGCGGGCAG  ATCACTTGAG
204181   GTCAGGAGTT  TGAGACCAGC  CTGGCCAACA  TGGTGAAACC  CTGTCTCCAC  TAAAAATACA
204241   AAAATTAGCA  GGGCGTAGCG  GCGCGTGCAC  CTATGCGCAT  GCATAGTGCG  CGTGCCAGCT
204301   ATTCAGAAGG  CTGAGGCAGG  AGAATTGCTT  GAACCCAGGA  CGTAGAGGTT  GCAGTAGTTG
204361   AGATCATACC  ACTGCACTCC  AGCCTAGGTG  ACAGAGTAAG  ACTCTGTCTC  AAAAAAATAA
204421   TAATAATAAA  AGAAAAGGAG  AACATGACCA  AAGTTATGAA  TAAGACTGAA  GGCAAGAAAA
204481   TTGTACGCTT  GTAGAGATCA  CCTAGCTTGT  TGCCCTCATT  GTACAGCTAA  GAAAAGGCAC
204541   CCAGGGACAT  TGTGGTCAGC  ACCAATTTCT  CAGAAAGATA  GGCAGATGAT  GAGAGGGCCC
204601   TCAGTTTTTC  TAACACTGAA  GGAATTGCTT  CTATGTTTTC  TGGTGAACTC  CTCCCCACTC
204661   ATCTTGAGGA  TTCCAGGCCA  GAAGAATCCA  CTTTAAAAAA  GAAACATTTA  AAACCAATTT
204721   AACAACCAAT  CAAAGGCACT  TTTATAGAAA  TACATTTCAT  TTGCTGTAGG  CCTGTATTTA
204781   TGGATCTGAG  AGGGCTAGAC  TGCCAATATT  GTGACTGTTT  ATTATTATTG  CTGTTGCTAG
204841   TATCTAGAAT  ATTATACAAC  ATATAACACT  TTGCAATTTA  CGAGGCATGT  CTCATACTTT
204901   TGTTTTCACT  CCAAACTGCC  CAGTGAAGTA  ACATTATCCC  AATTCTTCCT  ATGAAACAGT
204961   GAAAGCCCTA  AGAGTTTTTG  AAACTTTACC  TGGTTTACTC  AATTTGGGAA  TGGCAGAGCA
205021   GAATTCAGTC  CTTGAATATC  CTCCCACTGC  AGGTTCATGC  TCTTTGATCT  AGGTGTAACA
205081   TTTACTCTGA  GTAAACTAGG  ACTCTGGGCT  AACAGAGATG  AAGCAAGACA  GGCTGGATAT
205141   TAGGAGAATC  TAAGAGCAAT  CTAACGACCA  TTATAATAAA  ATCATGAGTT  CTAGACTTAA
205201   AAAAAGGGAA  AAACCTGTTT  TTTTGCTTAT  GCGTATACCA  TAATATTTAC  ATTATTTATT
205261   TTTTTCTCAA  ATTCAACCTA  TACTGTGTCA  AGTAATTTTT  TTTAATATAA  CATTTTCCTT
205321   TAACTTAATT  TCAATTCATT  TTTCTGTGTC  TACTTACAAC  TTTGGCACTA  GAATTCACAA
205381   TTTTTTTTTA  GAGGTATATC  TCCTTAAAGG  GAAGGGTTCT  GACACTGTTA  CATGTTCTCA
205441   ATTGTTTGCA  AATAGGTTAA  TAATTATTCC  AGTGTCTCTA  AGTACATATC  AACCATGCCA
205501   GTGTTCAGCC  TCCATAATTT  TATTAGCTTC  TGTGCTTATT  TTGGAAAAAC  ATTTCCCATT
205561   ACCATGAAAG  ACCTCAGTTT  AGGATGGTTT  GGTATGTTAG  CCTGATTTCT  GCATTCGTCT
205621   CATGCAAAGG  AAAATAGGAA  ACGAAGAACT  GAAATTACCT  ATTGATACAA  AATCAAAGTA
205681   GCATTTGAAA  CCATAAAACT  TAAGTAGGGC  TTTTCATCCT  TTCTCGTTAG  ACAGCAACAG
205741   AGAATGGGAA  GAAAAACTAA  AGTGATGGGT  TTGTGATACA  ATTCCAGTAA  CATAAAGAGC
205801   AAGGAGAAGT  AGTTTTGTTG  TGTTTATGTT  TAATATTCAA  AGCTCAACCT  AAAAGTATTT
205861   TTCATTATCA  AACTTCCTTC  TAGAATAAAT  GATTAAAACT  TGATTTAAAA  TATACAAATT
205921   CTCCTTTATA  ATACCTCAAA  ATGGAGCTAC  CCCATTGAGT  TTTAAGCTTG  TGATTAAAAT
205981   ATTACGAAAA  CAAAGGGGAA  GTTGTAATAG  GTAGAACAAG  CAGTAGTCTA  GGCATTAGGG
206041   GATCTGGTGC  TGGCTCTGTG  CATCATGTGG  TTTCAGGCAA  CTTTTCAAAT  TTTCTACGCA
206101   AATTTTCTTA  TCAATAAAAT  AAACAGTTGG  GCCAGAGGAT  CTCTGAGTCT  CTTTCAGCTT
206161   TCAGTGTTTA  TAAGATTGGA  GAAGTTGGTG  GGAAAGCTTT  AAGTGGAGTG  TAAGTAATTG
206221   CAGCTGCATG  TACAGTTAAA  GAGTTGCCTT  CAGCCAAGCC  ACGGGATCTT  GCATAAAAAG
206281   TGAAATCAAA  TAGAAAATGG  TCCAAACTCT  GGGTTTGACC  ACAGATGACT  TCAGCTAGGA
206341   TCTGAGTGTA  GAGCAATGAG  CTGAACTCCT  GATATCCAGA  TGTTAGCAAG  ACTTGGAGGC
206401   CTTCTAAGGC  AGAGCAACAA  CCAGTATCTG  TCCTGGTGCT  GACCTGATCT  TACTAGCAAT
206461   TGGGCCTCCA  TTTGGGTCCA  TTGTACAAAA  CAACAACAAC  AACAACAATA  AAATCTCCAA
206521   ACACCCAAAA  TTCAAAATTT  AGATGGAGAG  ATACTATTCC  CAGAATTCTA  GAGATATTTG
206581   GAAAGCAGAA  AACTATACTT  GCCATGCTGA  TGAAGTCCAA  TTATTGCTCT  TTTAAATACA
206641   TTTAGCTACT  TCTGAATATA  AAATGAGTAT  CTACTAATTA  TTTACAAAAT  CACTTGGTAA
206701   ATATAGAAAG  TCACAAAGAA  TGAAGTGATC  ATCCTGTTTT  GTAACCAGA   AATAGTCATT
206761   ACTGGCACTT  GTGTGAATCA  GTTTCTATTC  CTGTATGTGG  ATGTGCACAG  CGTATCCTGC
206821   TTTGTACACT  AGAGTACTAG  CATTTTTCTA  ATGTAATTCA  ATATTGTCGA  AAACATTTTA
206881   AAATAGCTTC  CATCACAATA  ATCTATCAAA  TTGACTTGCC  AGACTCTCAT  TATTAGGTTA
206941   ATTTATCTCT  AACATTATGC  AGTCATGAGT  AATACTACAA  AGGATATTTT  TGGACACAAT
207001   TTTTCATCTA  TGCCTTTCTT  TATAATCCTT  CATCCTAAGG  TCACAGATTA  TGAATATCTT
207061   TAAAGTACGG  ACAAGTCTTT  TAAATTTTGT  GTGCAAAAAC  AGTGCAAAGC  CTTGAATGAT
207121   AAAATAGAGG  TTTGATATAT  GTGTTTTTTT  GTTTGTTTGT  TTTGAGACGG  ATTCCTGCTC
207181   TGTCCCCCAA  GCTGTAGTGC  AGTGGCACGA  TCTTGGCTCA  CTGCAACCTT  TGCCTCTTGG
207241   GTTCAAGCAA  TTATCCTGCC  TCAGCCTCCT  TAGTAGCAGG  GTCTACAGGC  ATGTGCCACC
207301   ACACCCGGCT  GTTTTTGTAT  TTTTAGTAGA  GATGGGGTTT  CACCATGTTG  GCCAGGATGA
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207361   TCTCGAACAC CTGACCTCAA GTGATCCACC CACCTCAGTC TCCCAAAGTG CTGGGATTAC
207421   AGGTGTGAGC CACTGCACCC GGCCGATACA TGTGTTTTTA AAGTCACAGA AATTTCAGAT
207481   GTCTTGAAGG ATTTTAAGCA ATTTAAAAAA TAAAGTCATA GAAGCTTCAA TTTAGGAATG
207541   AATGGAAAAT TGATGATATT CTTAGGATAT GGATTTTTCC TAAAAGAAAC AAATGTATGC
207601   ATCCCCAAAG ATAATTTGAT TAGTATACAA ATATTAAATT AAACATGTCC ATATTTAGAG
207661   CCATGAATTC TCTTTGCCTG TCACAATAGC TGGATTTATT CACAATTGTA GTAATTAGTC
207721   CCTGTTCATT ATAATTTTCT AGGTGATATG AAGACTTTGT CAGTCCAAGC AAGTGTCCAC
207781   ATTGTGTGTA GCAAACATGA GAATAAACAT TTTAAACTTT TAAATGTAAT ACATATTAGT
207841   GTTATGTAAT GTCATCCTTC ATGTTCGAAG GCACATGGAA CATTGTTCTG GTGGTACAGA
207901   GGGGAGAGAA ACACCATCAG AATGAAAGGA AAGACCGCTC TGGAACCTTC CTCCTTAGCT
207961   CTTGAGCTTA GTTTAATTGT CCTGTCTTAT GGTCTGCTAC AAGCAATACC ACTCTTCACC
208021   TTCGCATGCT TCTCTGTGGT TTGATAAAGT ACATGCAATT TTTCATTTAA TTCTTCCAGC
208081   TGCACTAAGA AAGGAGCCTT ATCTTTATTG AACAGATGAG GAAATGAATG ATTAGAGAAT
208141   TTAAATGACT AGCTCTAGGT CACACAGCTG GAACTTACAG CCAGATTTCC TTTTAACAAT
208201   CCTGTAACCA AAAGCATACC AGTAGTGCCC CATAAAATGT AAGTTATAGA GCTGTGTTGG
208261   GTCAAAACTT TTACTGATGC TAAGAGGAGG CAACATTAAC AAGGGGAAAT TATTTGTGTA
208321   TTATGTTTTG GATTATGTTC TCTCCATAGA TAAAAGACTG TCGTAGTAAA AGAGATTCAG
208381   GGCACAGGGA AACTCCACCA CAAAGCGTGG TACCATTTCC CACAGAAGCT AAATGGACGG
208441   GAAGCCTGCC ACCAGGAAAG GTAAAGCCAC TGCTCTTGTT TGCAGGCTAT GTTAATAAGC
208501   TGAAGCTTAT TCCGACACAT TTACACATCT CTGCATCACA CTGACCCTTC GTAAAGATAC
208561   TCCCAGTGTA ACATTGGAGC CAGCTCCAGC CCCTGATCCT GTTGCTTTTT CCTTAGCCCC
208621   ATGAAATCAT CTGTGAGAAA TTAAGCCAAA TAAGCAATAA ATCCTGGGAT CTAGGGAGTG
208681   GAATAAGTTT TGGGAAAGTC TTTTTTTTTT TTTTTTTGA CTGAGTCTTG CTCTGTCTCA
208741   CAGGCTGGAG TGCAGTGGTG CGATCTCGGC TCACTGCAAC CTCTGCCTCC CGGGTTCAAG
208801   TGATTCTCCT GCCTCAGCCT CCCGAGTAGC TTGGACTACA GGCACACACC ACCATGCCCA
208861   GATGAATTTT TGTATTTTTA GTAGAGATGG AGTTTCGCCG TGTTAGCCAG GATGGTCTCG
208921   ATCTCCTGAC CTCGTGATCC ACCGGCCTCG GCCTCCCAAA GTGCTGGGAT TACAGGCATG
208981   GGCCACCACG CCTGGCCCGG GAAAGTCATT TTAAACCAAC CTATGTATGA ATCCCTACTA
209041   TAATATTCTC ACCAAGCGGC TGGCTCTTTC TCCTGAGCTT GGAAACCTCC AGTAAAATGG
209101   AAATAATTAT TTCCCAGACC ACCACTCTTA TCTGTGAGCT TTTTTGGCCA TTAAAAATTA
209161   TTTCTTCCAT TATATTTTTA TCTGTGTCTT CACAGGTTTT CTCTTTCTTT CACTTTAGTG
209221   CTTTTCTTCA AATAAGCAGG AAAAATCCAA TCTATCATGC ACATGGGAAC CCTTTCAATA
209281   TTGGTCTGTG GTTGTTCCAT TTTATGGGGA TGCTTTTAAA GAAAAAATTT GTCCTTTCAA
209341   TATATTGAAT ATCTTCCAGC ACCACATCAC CTGCAAGCTT TGTAAAAATA GTTCTACATA
209401   TTAATTTTTT TTTTTTTTT GAGATTGAGT CTCATTCTGT CACCCAGGCT GGAGTACAGT
209461   GACATGATCT TGGCTCATTG CAACCTCTGC CTCCTGGGTT CAAGTGATTC TCCTGACTCA
209521   GCCTCCCGAG TAGCTGGGAT TACAGGCATG CATCACCATG CCTGGGTAAT TTTTGTATTT
209581   TTAGTAGAGA TGGGGTTTCA CCATGTTGAC CAGGCTGGTC TCAAACTCCT GACCTCAAGT
209641   GATCCACCTG CCTTAGCCTC CCAAAATGCT GGGACTACAG GCGTGAGCCA CTGCACCCCA
209701   CGTAGTTTTT TTTTTTTTT AAGTTGAACA TATGTGAAGG CAGGACCTAG TGACACATAG
209761   CAATAACATT TCCAAGTAGA CATTACACTA GGGAATTAGT CGAAGTGCTC ATTTAAAGTA
209821   CCATCTCTCA AATGTATTAA AAGAGAATCC TTGGATGTGC AATACCTTAA TTCAAAGGCA
209881   GCTCGTTATG TATAAACTCT CAAGCTTTGT GATAAACAAA TGTGCATAAC AGATGGGACT
209941   ATTCACTTAC AGCCCAGGGA ATTTTATTGA CGCTGAGAAG GTTATGTGAC TGGCTCTGCC
210001   ACTGTCATCC CCATTCACTT CATTTGGAG CAATATGACA TAAATGCCTT ACATGTGGGT
210061   TTTCTCTATT TATCATGTGT TTCCTATCCC CTTGAAAGAT GGCCATATTT GCTTTACTTG
210121   GTTATAAGAT CCCATATTCG CTGTCTTGAA GCCAACCAAA TAATTTGACA AAGTGGGTTT
210181   GTAGTGCTGG CTATTTTGGT GAAAAAAAGA CAATGAGACT TCATGTGTCA TCCAAAGTTC
210241   TATCAGATCG AGCTGTGAGA GAAAGGAAAA GAAAGGGGTC TCAGTCAGGA TGCTCACTAC
210301   ATACATCTGT GTTGTTGTCT AGGTCCAGAT TTCTGTTCAT TACGCTATGG GCTGGCTCTT
210361   ATCATGCACT TCTCAAACTT CACCATGATA ACGCAGCGTG TGAGTCTGAG CATTGCGATC
210421   ATCGCCATGG TGAACACCAC TCAGCAGCAA GGTCTATCTA ATGCCTCCAC TGAGGGGCCT
210481   GTTGCAGATG CCTTCAATAA CTCCAGCATA TCCATCAAGG AATTTGATAC AAAGGTAAGT
210541   ATGATGGAAA ATAGGGCTCT TTGTTGAGAG AAAAAACTTT GAAAGGAAGG CATAGATCTT
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210601   GATTCTGTGG AGTATGGAAG TATACATTTC CAATGACAAA TTAAAACTGA CTGGAACTAT
210661   TTTTCTTTGA GACATTGCTT ACTTCAATAA TAAAAATAAG ATTTCATTGA GGTTATTATG
210721   ATTATAAGGT GGGGGAACTG TAGAGTTAAA TGTGAAAAAT TTAAAAATGG AACAGTTTAT
210781   GTGATGTCTT CAATGAAAAA CTAGGTATTA CCTGGGCACA TTCTTATAGG TTACTCAATC
210841   CTATTCAGTT CTCTGCCTGT TTTATTGTTT CTGAGCAATT TTATATCCCT GTAAATTCTA
210901   TATAACCAAT AGAAATGCAA ACGATTCTTG TCCATAGCTT TGCAAATAAA TTTTGCCAAG
210961   AGAAAAATCA GTTAAAACTT TTCTCCACTC ACCTCCCAGT TGAATTAGCC AATTTTGCTG
211021   TTTGTTTGTT TGTTTGTTTT TTGAGATAGA GTCTTCCTCT GTCATTCAGG CTGGAGTGCA
211081   GTGGCATGAT CTCAGCTCAC TGCAGCCTCC GCCTCCCGGG TTCAAGAGAT TTTCCTGTCT
211141   CGGCCTCCCA AGTAGCTGGG AGTAAGGGGG CATGCCACCG CGGCTGGCTA ATTTTTGTAT
211201   TTTTAGTAGA GACAGGGTTT CACTAGGCTG GTCTCGAACT CCTGACCTCA GGTGATCCAC
211261   CCGCCTCGGC CTCCCAAAGT GTTGGGATTA CAGGTGTGAG CCACTGTGCC AGGCTCTGCT
211321   GTATATTTAA AGTCTATTTC AGCATTGCTT CCTGCTTGTG TTATGCGTGA TTCTTTGAGT
211381   TTTCCTTTGA ACCAGTTATA ACATCTTACT TACTTCCTCC ATTAATCAAT GAGTTAAATA
211441   AAATCTTGT  TGTATGTTTA TTTTACATTT ATATGAAAAC CATGAATTTA CCCAATTAAA
211501   AAAATTATCC TTTAAATTAT CTTGTACTGT ACATTCCCA  TGTCATCCCT ATAATTCATG
211561   ATTAATGATT TTATTACATT GGACCTAGCT TATTTACAAT GAGTACATAA ATTTATTGTC
211621   TCCAGTCTTT CCTCCATTAT CCCGTCTACA TATCCACACT GAGTAGATTC ACTACTCAGG
211681   AATCTTGGAC ACCTTCAAGT TGCCAAACAT GCAGTGTTCA CTGGACATGC TGTGTTCCTT
211741   CAGAATTTGG GCCTGCTTCT CAGCACACTC ACATCTGCTA TCAATGACCC ATGGAAAGTT
211801   TTTGCCCTGA GCAAGCCAGA GTCCCTGTTA GTTTCTTCCA AATGCTACAA GTTCACTTTT
211861   GCTATTTTTT CCGATGAGAT AAAATTTTCC TTTTTGACTT TCTACAAATC ATAGTCATTT
211921   TTCAAGGGAT AGTTCAAGTA TTGCTTCCTT TCTGGGACCT TCCCAAATTA TTATTTTCTC
211981   CTCTCAAAGT CTCTGTTTTA TTTATGTTCA TCCTCAAATC TTGATTCTCA CATGAATCAT
212041   ATACCTTGTA TTATTTATAG TTTTTTTGAG TGGGTAAAAT ATTTCATATT TTATATTCTT
212101   TGGCTCTCTA CTTTATAGCA TGATGCCAGA TATTTAGGGG CCTTATTGCA TTTATTTTTT
212161   ATTTTATTTT AAAATCTATT TTATTTTTTA TTTATTTATT TTAAAATCTA TTTATTTTTA
212221   GGTAAATATT CAGGTAATAT AATTTATGTA ATTATTTAGG AATTTTAGGT AGTTATTTTA
212281   AAATAATTCA AATTATTTAT TGAGTTATAT CAGAAGAATG TGATCTTATT CATTTGTAAT
212341   ATGTGTTTTA GGAACTCAGT TCAGCCAGGG CAGACCATGA TTCCCAAACT TGACTTTTCT
212401   TTTTAATTAG GCACTGATTT TGGTTAAGAG TTCAGTAAAG TTTTGTGTGT GTGTTTAAA
212461   AAATTCTTTG ATATAAGAGT CAAGATGTTA CTCAACTTTT ACTAGAAGCA AAATAGAGGA
212521   AGTGCTTTCA CAGATGAAAT ATCTCTCAAT GTTTTCTTCC ATTTACTTCT TCCTATTATT
212581   CATCTATATA ATCATTTTCT TTACCTCTTT TCTTCATTTC TTCTGTTTTT CTCTCCTTCT
212641   ACTAAGACAA GCAAATTAGG GGTATAATTG GTTATTTGGG AAGGTAGGAA GAATATAGAG
212701   AGAAACAAAA ATCAATATTT TATACTAGGG TCTCACTAAC CTCAAGCAAC TCTGACTGTA
212761   AAGTAGATTT TCATAATAGG ACTTCTTGAC AAAGAGTTTT CCTATTTTTC CCCCAGGCCT
212821   CTGTGTATCA ATGGAGCCCA GAAACTCAGG GTATCATCTT TAGCTCCATC AACTATGGGA
212881   TAATACTGAC TCTGATCCCA AGTGGATATT TAGCAGGGAT ATTTGGAGCA AAAAAAATGC
212941   TTGGTGCTGG TTTGCTGATC TCTTCCCTTC TCACCCTCTT TACACCACTG GCTGCTGACT
213001   TCGGAGTGAT TTTGGTCATC ATGGTTCGGA CAGTCCAGGG CTTGGCCCAG GTATCCAGAT
213061   ACTTTCTCAT TCTTGGTGGG ATCCAGATTT CTGAATTCTA CAAAATATCA AAGGTCTTAA
213121   TGATTTTCAT TTCAGGGAAT GGCATGGACA GGTCAGTTTA CTATTTGGGC AAAGTGGGCT
213181   CCTCCACTTG AACGAAGCAA GCTCACCACC ATTGCAGGAT CAGGTAAGTG TGCACAGATG
213241   GGTCATAGCT TTGTCATCTG TTCCATCCCA CTGTGTCTTA TCTTCTATGA ATCAAATGGT
213301   TTGGGGAAGA GAGAGAAAAA GTACTGCTGA AAAATTCAAC AATATAAGAC ACTTGCATCA
213361   CAAATAGGAA AGATGCATCT GTGCAGTAAA GACATTGAAG CTTAGAAGTA GAAAAAACCA
213421   TTGTGAGCTA GGTTTCAGCT CAGAAAAGCC TTAGTAGTCA GAAAAGCCTT AGTAGTCAGA
213481   AAAGCCTTGT CGGAAAAAGT TTAAACCTTT AAGAATTGCA CACATGGAAA AAGATCAAGT
213541   AAGCTATATA TACACCATCT TAGCAATGAT TTTGAAGTGA GAATTAAGGC TACCACAGCT
213601   CCAGGTGGTA AGGAGAGAAA TCAGGCTGGA AGAGTTTGAA GTTTCTGTAT TATTCTAAGC
213661   TCTTTACTAT TCTATTATGA GCTCATTAAT TCTCACAACA ACCCTCTCAT ATAAGTACCA
213721   TTTTAAATTC TTATTTTACA GAGAAGGGAG TTAAGGAAGG TGGAGATTAA GAAAATTGCC
213781   CAAATACAAA TAGCCAGCAG GTGGTAGGTC TGAGATTTAA GCCCATGCAG ATTTTAGCCC
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213841  CAGAGCAGAC ATTCTCAATC ACTATGCTAG ACTGCCTTTC CATGGTATGT GATCCTACTC
213901  AGGCCTCTAC AGCTTTATCA TTGCTGTTCT CCCCAGCCTG TCGTGCTGAG AGTATATACT
213961  CGAAGAGCAG AACTAAAATT CCATCCAGCT TCTCACTCCT AGGTCCACTA CACAGCTGCA
214021  TCCTGCAGAC TTTTACCTCA AGCAACCCTC CTGCGTTCTT GCTTCCTTCC ATCATAGTTG
214081  TAACCATCTC CTCTATTTGC AAATACTATC TGCTGATCTC TCTCTTCTAG ACTGGTTTCT
214141  TTCAACCTTC TTCCCACCAA AACCAAGTTA GCTTGCTAAA ATAAAGATGG CACATTTTTA
214201  CTCACCCGCT TGAGAATTTT CAATGTGTTC CTTCATGCTT ACAGAGTAAA GCCTGACCTC
214261  TTTATTGCAT GAATACAAAA GTTCTTAGCC ATCTGGCCCC AACCTTGTTC CACTCAACTC
214321  CCCTGTGCAA GCATGGCTCC AGTGGCACTG GACATTGGCT GCTCTCCACA TAGATCTGCA
214381  CTGCACTTCC CTCTGGCTCT GCTCCCGTTA GTTTATATGC CTGGAAAGTT CTTTGCCCCT
214441  GTTCCTTGTG CCAAAATTCC ATCTATCCTA TTGCATAGCT TATGTAAAAA CTTCCTAAAC
214501  CTTTTTTTTT TTTTTTTTT TTTTTTTTG AGACGGTGTC TCACTCTTTC GCCCAGGCCG
214561  GACTGCAGTA GCGCTATCTC GGCTCACTGC AAGCTCCGCC TCCCGGGTTC ACGCCATTTT
214621  CCTGCCTCAG CCTCCCGAGT AGCTGGGACT ACAGGCGCCT GCCACCATGA CCGGCTAATT
214681  TTTTGTATTT TTAGTAGAGA CGGGGTTTCA AGCCAGGATG GTCTCAATCT CCTGACCTCG
214741  TGATCCGCCC GCCTCGGCCT CCCAAAGTGC TGGGATTACA GGCGTGAGCC ACCGCGCCCG
214801  GCCAAAACTT CCTAAATCTT ATAATTATTA TCAATTTATC CTCAGATATA CTTCCACGTA
214861  CATTGTAGTT TTATTATATT TATATTTTAC ATCTTTTTTT TCAAATTTCA GTTTGGGACC
214921  CATTAGTGAG TCATAAAATC CATTGAGCGG GTTAAAATCA TTATTTAAA AAATGAATAG
214981  AATAGAATAG AAATTGTTGG AGTGCATTGG ACATGGTAAA GTTAAATATC GATTCATGAA
215041  ACCATCGTTT GAGGCATATG TGTGTGGTTG TATGTACAAG TGTTATGCA TATTGGTGTG
215101  TGTGTTATGT TACCCTGTAA AATGCATTTC TTACTATAGG TCTCTGTGAA ATATGTGTCT
215161  TGTTGTTTTT TAATGTAGAC TTCCAAAGCC TACATGGCAT TTCACTAGTG ACAATCAATT
215221  TTATTCACAT TTTTCTCTCC AATTGGACCA GAAGCTCTTT GAGGGCAGGG GCTGTATCTT
215281  ACCGATTTTT GTAAGTCTTT CATTTCCTGC CCCTAGCCTC ATATTAGATC ATGCAAGAAT
215341  GCAACTGTAA TCACAAGAAA ATGCTAATGG GCTGTGATAG CAGAGAGTTA CTGTGACAAA
215401  CTAAGGGATT TAGATTTGGT CACATTGGTG TTGAGGAGCC ATTGAAGAAT CAGAGAGTGT
215461  GTTACTATTA TTTGTTAATT TTAATTATAT CATATTACTT TACTGGGGAA AATCTGTGAG
215521  CTATTTTAGA AATAAATACT CTCATTGCCC AATAATTCTA AGTCTGCCAC CTCACTGTTG
215581  GGACATTGTT TAGGGAGGCC ACGAAGTCTC AGCCTTTGAT ATTTTCATAA GTGTTTTTCT
215641  CCCTTTTTCC TTTAGGGTCA GCATTGGAT CCTTCATCAT CCTCTGTGTG GGGGGACTAA
215701  TCTCACAGGC CTTGAGCTGG CCTTTTATCT TCTACATCTT TGGTGAGTCA CTTTCTCTTA
215761  AATCCTAACG CCTCCATTTC CTGAGCATCC ATTTTGGCAC CTACACCACC CACATTCTTC
215821  CTATATGAAA GAAAATGTCC TTTATCAAAT GGAAGATGAT AAAAAATGTC AACGGTTGGT
215881  ATCATTTTTA ATCTAGTCAC ACAACCTGAT TAACACCTTC CTGGTGGTTC TGGGAAGCCA
215941  CACGCACAAG GTAGAGGAGT TGACTATTCA CATGGCACCC ACCGACTTGT GATGCAGTCT
216001  TGTCCTTCCA TATCAAGCAC CTTCTGCAGA ATCTCTACCA CCACATCTGA AGTGCCTGCT
216061  ATATGCAGTT AAGATGTCAA AGATAGTGAA GTACATTTTC AATGTGTCTT CATATTTCAT
216121  TATAATTATT ATTTCTGTCC AAGATGCCTT TCACCTGTTC TCTACCAAGT TAATCTTGCA
216181  AAGTTCAATT CAAATGTTCC CTTCCCCATG GGCCCTTCCA GGGCTTACCC TATCAGATTC
216241  TGGCATTCTC TCCTTTATGA TATTTCCTCT CTAGGTTATG TTGGTGTGTA ATTATTTATT
216301  TCTCCTTTTC TTTCCACTAG ACTGTGAAAT GCTTGAGGCA AGGAATCCAT TCTATGTTTT
216361  CATCACTTGG GTGTCATCAT GGTGCCTGAT TTTTAGCTTT AAAATAAAAG AATCAGTGAA
216421  TCCAGTAATT AGAGGGGATT TAAAGAAAAC TAGTCCTCAG AATCTTTTAA CATAGAATGT
216481  TCTTCAAATA AGGAATTCCA ATAATAAGAC AATTTCTAC ACTTGATTTT GTTTTTATAG
216541  CCAAATGGTG TCATTAAATA TAGTCCTGGC CTGAATGGCT TTCTCATTAA TGATGCTAAT
216601  TATTTTGGTT TGTACATGTT AACCAGGTAT TGTACAAAAA TATTTCTTTT GGGAATCCAT
216661  AATGGATGTA TGGCTTGAAT ACAAATAATA CTGTCTCTTG TAAGTGCATT GGAAATTTTT
216721  CCCTGCCACA TGATTTCATG GAAGGTTGTT TCGTGTATGT ATGACTGCAA ACCTGACTAT
216781  TCAGATCTTC CGCAACAAGA CAACTTATGT GTGCATTAAG AAGTTGCTGC CTAAAATACA
216841  TAACACTGTA ATCATTGGAG ACTTTAAAGT AATTAATCAG CTATGCAATG CCACGCTCCT
216901  GTTATCTCCA GAGGGCTCTG ACATTGACAA ATGGTGGCTT TCTATTTGAG ACGTAATATC
216961  TAAAAAGCTT TAACAGGTTT GTAGAAGGAT TGAAAGAAAG AATGGGAACA TTTAGGTCCT
217021  TATGGTAGAA TAAGCATTAA TTGATTAGTG TGTAGAAGGG AGAGGCATGC CACTTCAGAG
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217081  GAAACTTCCT  TCCCCCAGTA  AACAAATCTA  CCTAAAAACT  AATTTTATCC  CTTCTTCCCA
217141  GGTAGCACTG  GCTGTGTCTG  CTGTCTCCTA  TGGTTCACAG  TGATTTATGA  TGACCCCATG
217201  CATCACCCGT  GCATAAGTGT  TAGGGAAAAG  GAGCACATCC  TGTCCTCACT  GGCTCAACAG
217261  GTACAGTGCA  CACCTTGTAC  CTGTGGCCCA  TGCAGAGGTC  TCTAGGGCAG  GGTGTGGATC
217321  TCCTCTGAGA  GGCACCATCT  TGGCTGCTCT  AATACTCATG  CTGATTAGAT  CTTTCTTTTC
217381  AGCCCAGTTC  TCCTGGACGA  GCTGTCCCCA  TAAAGGCGAT  GGTCACATGC  CTACCACTTT
217441  GGGCCATTTT  CCTGGGTTTT  TTCAGCCATT  TCTGGTTGTG  CACCATCATC  CTAACATACC
217501  TACCAACGTA  TATCAGTACT  CTGCTCCATG  TTAACATCAG  AGATGTGAGT  TTACTTCCTA
217561  TACTTCTACG  AAAATGATAA  TGGTAATAAG  GAGAAACAGT  TCTGTGTTAC  CTATTACATT
217621  CTGGCTTTAC  ATATAACCAT  TAATTTAACC  TTCACAATGA  CCTTGAGAGA  GGCATTGTTA
217681  TAATTCCCTT  TTCACAGATG  TGGAAACAGG  ACACTTAGAG  GTGAGATAAC  TTGCCCCAGG
217741  TTGCACAATA  CTAAGTGATA  GAGCTGCTGC  AGCATCCATA  TTCTTAACCA  CTATGCTATA
217801  CTACCACACC  AGCTGATTCC  AAAGCTTCTT  TTAGAAATAA  TATTGCTGGG  CCAGGCATGG
217861  TGGCTCATGC  CTGTAATTCC  AGCACTTTGG  GAGGCCGAGG  CAGGCAGATC  ATGAGGTCAG
217921  GAATGCAAGA  CCAGCCTGAC  CAATATGGTT  TACTAAATAT  CATCTACTAA  AAATACAAAA
217981  ATTAGCCAGG  TGTGGTGGCA  GGCACCTGTA  ATCCCAGCTA  TTCAGGAGGC  TGAGACAGGA
218041  GAATCGCTTG  AACCCAGGAG  GTGGAGGTTG  CATTGAGCCA  AGATCATGCC  ACTGCACTCC
218101  AGCCTGGGCG  ACAGAGTAAG  ACTCCGTTTC  AAAAACAAAA  AACCCAAGAA  ATTAATATTG
218161  CTTTTATCTG  GAGCCCAGAG  TGATGCAGCT  TCTGGCCCTC  TTATCTGAGA  CAGTGTTCTT
218221  TTAGTGTGAA  AAAGGATGCT  AATTTTCCCC  CAAACAACCC  ACAGTATCAT  GGGGGTAAGT
218281  TAATGGCTGG  TCTGTGTAAC  TGACAAATTT  TGGTGCTAAC  GTATCTCTAT  AACTACTCTG
218341  TATAAACTTC  CTTCCTTCAG  AGTGGAGTTC  TGTCCTCCCT  GCCTTTTATT  GCTGCTGCAA
218401  GCTGTACAAT  TTTAGGAGGT  CAGCTGGCAG  ATTTCCTTTT  GTCCAGGAAT  CTTCTCAGAT
218461  TGATCACTGT  GCGAAAGCTC  TTTTCATCTC  TTGGTAAGGA  TAAGCGTGTG  GGCCCATTTA
218521  ACCAATCCCT  TTTCTGCACA  TGGTCTCAGA  GGGTTCCCTG  ACAGCATGTC  CTCATTGCCC
218581  AGGGCTCCTC  CTTCCATCAA  TATGTGCTGT  GGCCCTGCCC  TTTGTGGCCT  CCAGTTACGT
218641  GATAACCATT  ATTTTGCTGA  TACTTATTCC  TGGGACCAGT  AACCTATGTG  ACTCAGGGTT
218701  TATCATCAAC  ACCTTAGATA  TCGCCCCCAG  GTAAGAGCTC  TACCTGTTTT  TTCCCCTCCT
218761  CCAGACCCCT  CCAGAGGTGT  TAGACCTCAG  TGGTCGCCGT  GAAACTCTTT  AATGTTACTG
218821  ACATTGCACT  AATGGCAGAA  TGACAAATAA  CTACAAATAT  CTGTCTGTGG  CCATTTTTAG
218881  AACAACAAAT  GTGGCATTTT  TAGAACAACA  ATTTCCAATC  TTGGCCAGTA  ATCATTTGA
218941  CAAAAACCTT  CCCAAGCTTC  CCTAACAGAG  ATTGAACTGT  GTATGCTGGG  AAAAGGCCCA
219001  CACACAGGTG  ATTTGGAAAA  GTTTCCATGG  TGTTGTTCAT  ATTAGCTACC  ATATATATAT
219061  ATATATATAT  ATATATATAT  ATACAGTCAC  AATAAGCCAG  CTCCTGTGCC  AAGACTTGCC
219121  ATATATCAAC  ACATCTAATC  CTCACAGTTA  TATTAGGTAG  GCCCTATTGT  TATCCCCATT
219181  TTATAAGGGA  GAAGGCTGAG  GCACAAGGAG  GTTAAATGGT  GTGACTATGG  TCACATAAAG
219241  GCAGAGCCAG  GATTTGGACT  GGGGGAGTCT  GGCTTTGGAG  TCTGTGTCCT  GCCCGTTGCA
219301  CAAACTGGCT  TCTCCACTGA  GCAGCCGGGG  TAAAGAAACG  TGGTTCCCAG  AGAGACTGCA
219361  TTGCTCCCTG  GTTATTGACT  TGGTAGATTG  GTAATTTCAG  GTTTGGCAAA  TAGACATTGC
219421  CCTGAATGTC  TTTAGGTGAA  TGAAAACTG   CATTAAGCAA  AATGACTTTG  CCATTAGAGC
219481  TGAATTGCAT  TAAAGTTGAG  TTGCTGCAGA  AGCTGTAGGT  GGCTTTCTAT  ATAAAATCAT
219541  TTATAAAATC  ATCTTCCCAC  AGATATGCAA  GTTTCCTCAT  GGGAATCTCA  AGGGGATTTG
219601  GGCTCATCGC  AGGAATCATC  TCTTCCACTG  CCACTGGATT  CCTCATCAGT  CAGGTTGGGC
219661  CAGTTTATTG  AACATCTTCA  AGTGGCAGGT  ATTGTTTTAG  GTGTTGGAGA  TACACACGGT
219721  GCTCTAAAGA  TCTGGATGGC  AACACAATTA  CTCTATTTAC  ATGAGCCTCT  AAATCAGACT
219781  CTGGTAGGTC  AGATTTCCCA  GAGGAAGAAA  AATATAAGCT  TATTTTCTCA  AGATGAATAG
219841  ATGTTAGATT  GATTAAAATG  AGCTGTTCCG  GTGCAGAAGA  CAGCACGTGT  GACTTCCTAG
219901  AGGTACATGA  GCATGAAACA  GTTCTTAGTT  ATGACCAGAA  TGAAAGACAC  ATGTCAAGGA
219961  ATAGCAAGAG  ACGAAGACAG  AGGGGCAAAA  GAAGATCATG  AAGAATATGT  TCAGACTAAT
220021  CCAATTTTTA  AAAAATCACA  AAAGGGAAAC  AAAGTGTCCT  AGGCCAGTTT  AAAGATAATT
220081  TAATGTCTGG  AAACAGATCG  GCTGTGAGAC  ATTGCAAGGA  GGCTTGCTCG  GTGTTTGGAA
220141  ATGCAGGCTC  ATGAGGAAGA  TGAAAGACA   GACCCAGGCA  GGGATGGAAG  GACTGACGAG
220201  AACCAACTTA  CAAAGAGAAG  TTTTGTTTTT  ACTACATTTC  TATGTGATCA  AGTTCCCAGG
220261  TTAATATTTG  ACTAAACTGC  TAGGAATCCA  CTGTGACTAT  AATGCTGGAA  ATGACTTAGT
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220321   AGGGCTTTCT GAGGAGGGTC ACACAGAAGA CCAAAGAGAA CTCATGTTGA ATTGAGATGG
220381   GTTGTAGTGA TAGTTGTCAA CAGCCAATAC AGAAACAAAA AAAAACAAAA CAAACAGCAA
220441   CAACAACAAC AAAAAAAAAC AGAGAAGACA CAAACACAAT GCCACAATGC CATTTTAGGC
220501   ATAATTTTAA ATGAGTAATA TTATATGTTG AAATCCAAAT TTTCAGAAAA ACATTAGTGT
220561   ATTTTATTTT TGTTTAAAGA AATAACCATC TCAACTCAGA ACCCCATGTG CATTTTGGCC
220621   ATTTTGTTTC CAATAGTTTC ATAAACTTTC TTAAGTAACT ACTGCACATT GTTCCTTATA
220681   TTCCTTGTGA TCAACATTGC AATACACAAC TGGGAGGGCT ACTAGAACTG GTGTAGAAGG
220741   AACTTGTGAG ATTGATCATT TTCTCTGTTT TTTACATCTA GGATTTTGAG TCTGGTTGGA
220801   GGAATGTCTT TTTCCTGTCT GCTGCAGTCA ACATGTTTGG CCTGGTCTTT TACCTCACGT
220861   TTGGACAAGC AGAACTTCAA GACTGGGCCA AAGAGAGGAC CCTTACCCGC CTCTGAGGAC
220921   ATAAAGTTAC AAACTTAAAT GTGGTACTGA GCATGAACTT TTTAAACATT TTTTACTTCT
220981   CTCCATATTC CTGACCATAG ACTCAGCAGT TCTTAACTCT GGCTGTGTGT TAGTCTTCCC
221041   TGGGGAGCCT TTATAAGACA CTGATACTTG GGACCCACTC CAGAGATTCT GAATGAATTG
221101   GTCTGGGGTG GAACCCAGAT ACTACTAATT TTTAGATACT CCTTAGAGGT TTCTAGCATG
221161   CGCCCGGGGT TGACAACAGC TGGACAAACT TGAAAAGTCA ATTCATGTGG CCTTTGAATT
221221   TTCCTCATTG GAAAGTACTA AATAAATAAA AATTCATGTG AAAATGATCA CTGATAAATA
221281   TCTTCATGGT GGGGCAGGTT ATTGGATGCA GAGAAGATCT GCTCGGAATT GTAGCCATAT
221341   GTTACAGATC TCAGCACCGA TCGGAACTGT AAAGCTATAA TCCCCAGAAT TAAAGTTTTT
221401   ATTATTTTTT ATACATTGTA AAACATAGAC GTTTATTTAT GTGATTAAAT TCTATTAAAA
221461   TTTACATGCT AAAATAAAAT AGACCATTTT CAAATTATTT AGATCCAGAT ATTTCCATCA
221521   GATTAAACAG ATATTTATTT ATCCTAGCCC AATTGCAAGA GATTAATGAT GAGAAAATGA
221581   CCAATACAAG ATTAAATAAA TGAGGTTAAC TTAGAAATCA AGGACAGAGA AGATAGAACT
221641   GGAAGGCTTG TATTGTGAGA AGAATGAATG TGAAGGAAGG CAATGTAGAC ACTTCCAGAA
221701   GGGATAGCAA TATAGTTTAG ACCATATAAT GAAAATTGGA GAGAGATGAC AGAGACACTT
221761   TCAAGTGAAA TGACAATTTA TATGGGGGAG AAAAATATTG AAGACATAAC AAGATGAGAA
221821   AAGGCATAGA AATGTATCAC ATACAAGGCA TAGAAGTGTA TCACATACAA GAGAAGTTCC
221881   TTTTGAGCGT AGAAAAAGAT AATTTAACCT TCTTCATATT TTTCTTACTT TCCCAAGATA
221941   CTCAGATAGG CAGCGTCAAC TCTAACAGGA ATTAATTTGG CTCCTAACAC TTAAGACATA
222001   TCCTTTAGTT TGTCTCCTCA CACAGAACTG ATTCTGGTTT TGCCACAACA TGTCTAGAGA
222061   AGAAGTTCCC ACCATATTTT AAATCCTATT AAAAAACTGC TTGGACAAGA ACCTTGGGTT
222121   AATTCAGCAG ATGAAGAGAA TCTCCTAATG CAAATCAATG GGTATTTTTG AGCAAGTTTT
222181   TCAGAAAAAC AGAGTGTCAG GCCCTGAGGG TGGTACTAAG ATGAGAACAT TGATTTTGCC
222241   TTCATGATAT TGACAACACA AAGAGGAAAG GGGGTTTGCA GAAAACTAAA AGAAGAAGTA
222301   GAAGAAAAAA GAAAGACATA GTATAATAGG TAGTCAAATT ATGTACAGAA AAAAGAGAAA
222361   AAAAAAACAA AAAAGGGTGG GGGACAGACA ACCCAACTAA AAAATGGGCC AATGACTTGA
222421   ACAGGGACTT CATAAAAGAG AAAATGTAAG TGGCTCCTTA ACATATAAAA AGATGTTCAA
222481   CTTCATTAGT CATTACAGAA ATGAAAATCA AAACTACAAT GAAATACCAC TATAAAATTA
222541   ACTAATGGAT AAAATGAAAG GAGATGGAAA ACAAAATGTT GCCAGACATG TGGAGCAACT
222601   GGAACTTTCA TACGTTACGA ATGTGAACTT TGGAAAGCTG CTCGGCAATA TCTCCTAAAG
222661   CTAAATGTAC AATTCCAGTG ACTCAAACAT TTTACTTAGA AATGCACATA TACATCCATA
222721   AAACATGTAC AACAATGTTC ATAGGAGCAC TATCTGTAAT AGCCTGAACA GGAAGTTGTC
222781   TGTTAAAAAA AGAATGAGTA AATAAACCAC GGTCTATTTG TATAGCAATG AGAATTAACA
222841   GACCCCAATA TATAATAGAT GAATGGGTCT CATAAGCACA ATATTGATTA AAGGAAGACA
222901   AAACGCACAT TCTTTTAAAG GTTTATAAAA TACTTTTTAA AAACAGCTAC AACCAATCTG
222961   TCCTGTTAAA AATCAGTGAG CGATTCCCT TGTGCAGGGA TGGGGGTTGT GGCTGGATGG
223021   ATGGTACTTA AGAAGTGCTC CTGGGGTACT AGAAATATTT TATTTCTTGA CTTGGATGTG
223081   TGTTACTTT GTGAATATTG TACATTTATG ATTTGTGCAC GTTTATGAAT GTAGAAAATA
223141   AAACAGAAAG CAAATTCAAA GTATCATCCT TTTGAGAGCT TCTGCTCTGA CTTCGTTTTG
223201   ACCAATGGAG CAGTTGGGAA GGGGTCTTGG TCCTTCGGTC CTTTGCTTTT TTTTTTTTTT
223261   TTTTTTTTT TAGACAGAGT CTTACTCTGT CGCCCGGGCT GGAGTGCAGT GGCTCGATCT
223321   TAGCTCACTG AAAGCTTTGC CTCCCGGGTT CATGCCATTC TCCTGCCTCA GCCTCCCCAG
223381   TAGCTGGGAC TACAGGCACC TGCCACCATG CCCGGCTAAT TTTTTGTATT TTTTAGTAGA
223441   GACGGGGTTT CACCATGTTA GCCAGGATGG TCTCGATCTC CTGACCTCGT GATCCGCCCA
223501   CCTGAGCCTC CCAAAGTGCT GGGATTACAG GTGTGAGCCA CCGCGCCCGG CCCCTGGTCC
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223561  TCTGCTTTCA TGTTCTTCTT GGTCCTGTTC CTCCTCCTCT TTTGTTGGAA CTTCCAGTAT
223621  CAGAGCAGGA AGGAAGGCAA TGGGTCAATC GATGCTGTCA GCTTTTGGAT CAAACTGCAA
223681  GTTCTCAAAC AGCAAAATTA ATGAGCTCAG GCTTTGAAGA AACCATGACC CTGAAAGCAT
223741  CAGTTGCTTC CAATTGCATC AGTTGCCACG GGTGATAAGA ACAATGATGA CTCAGAATGC
223801  CTAGGTTTTC CCAGCAGCTT CTCTGAGGTT TTCCCAGCAG CTTCTCTGAT TGATTCCTGA
223861  CAGATGACTT CGGTGTGTCA GACTTTCAGG GTATCTTTCC TTATGTGATG GTTTGAGGAA
223921  GAGTTACCAT TCACATTCCT AATGGCTTCA GAATAGATGC AATTGTGAAC TGATAGGAAA
223981  CATTTCTAAT TCATCTCCCC TCCCCATCCC TAAAGGATTG TTTCTAACAA TAGTCATGAA
224041  AATTAATTCA CTTTTCTCAA ATAGTTTATT GTCATCTACC TAATGATGAG ATGACTTACT
224101  TTTTCTCCTT GACTGTTAAA TATTATGAAT TATATTAATG TATTTCTTAA TGTTGAGCTT
224161  TCCCTTGAAT ATTCTTTTGA TGTACGACAG AATTTGATTC ACTAATAGTT TATTTAGGAC
224221  TTTGGCTGAT GTACTGATAT ATGAGATTGG CTCTGTATGC ATACATGTGT TTTGTGTATC
224281  TTTTTTGTGT CTGGATATGG AGCTTATGCT GATTTCAAAA ACAAGAAAGG AGAACTTTCC
224341  TTTTTCCCCA TTACTCTGAA AAAGATTGAC TAGAATGGAA TTTTTATAAT TGCTGTTGTT
224401  ATTTGAAAGC TTGAAAGCAT TGGTTTGTAA AAATCATGCA GGCTGAAAGC CATTTTGAGG
224461  AGACTTTGAT AACTTTCTCA ATTTCCTTCA GTTACTGGTC TTTTAAGGGG TTTTATATTT
224521  TTCTTTGATC AATTTTGACC ATTTATGTTA TCTTGGAGGA TCATCTATTT TACACACTAT
224581  TTAAAGTATA TTTGCAAAAA TTCAACTGTT TTATCAGGCT ATCTTTTTAA TAATATATTC
224641  ATTTTATCTA TATCTGAGGT TTTAGCTTCT TTGTACTTCT GACCCAATTG CATGTGTGCT
224701  TTCTTTCTCC TTCATTAGAC TACTTAGTCA TTTACTAATT TTAAGAATAG CTTGTCTTTT
224761  ATTTATTTAC TTATTTATTT TTGAGACGGA GTCTCACTCT GTCACCCAGG CTGGAGTGCA
224821  GTGGCGCGAT CTCGGCTCAC TGCAACCTCC GCCTCCCGGG TTCAAGTGAT TCTCCTGCCT
224881  CAGACTCCCG AGTAGCTGGG ATTACAGTCA TGCACCACCA TGTCTGGCTA ATTTCTGTAT
224941  TTTTAATAGA GATGGGGTTT TGCTATGTTG GCCAAGCTGG TCTCAAACTC CTGACCTTAG
225001  ATGATCTACC CACCTTGGCC TCCCAAAGTG CTGGGATTAC AGGCATGAGC CACTGCGCCC
225061  AGCCCTGCTT GTCTTTTTAT TTTATATTTG ATTAGCTTTA TCTTTTATCA AGCTTATGTC
225121  CTATTTCCCT TTGCTTTACT TCATATAAAT TTTGTTTTGG ATAGTTTATT TATTTTTCAT
225181  TTAATTATGA AACAGGTTAA AGCTTAGAGG AAAATTGCTC CTCTAAGTCC AATTTTGTGG
225241  GCAGATTACA TTTTGCTGTG TTGTGCTCCC AAATTCATTG TTCTTTTAAT GCTTTATTTC
225301  TCAAGTTAAT AACCTATATA GTAAAAAAGT GGCTGTTGAC TCTCAGCTTT TTTTTTTTTT
225361  TTTTTTTTT GTAGATACAG GGATCTTGCT GTGTTGCTCA GGCTGGTCTG AAACTGCTGG
225421  CTTCAAGGGA TCCTCCTGCC TTGGTCTCAC AAAATGCTGG GATGACAGAC ATGAGACACC
225481  ATGCCTAGCC ATGTCTCTCT CCTTATATAT AATAAGAAAA CAGACACACT GAGGCATCCT
225541  ATCATCTCAC TCTTGGTTTC ACTACTGTTC TCTGGAAGTT TTGCTCTGAC CTTTTGCAGT
225601  TAATGTATTA ATTTTGCATT GAGTAGTTTC CATAGAAGAA TTATAGCATT TGCATTCTGT
225661  TGGGTATTAT ACTTTTCACT GTTATTTGAA CATAATTTGA GGGCTGAAAC CAAGATGAGG
225721  CAAGTGAGGT GCCCAGGAAG CAATATTTAA GGAGGCATCC TTTCTTAGGC TCATGCAAGA
225781  ACAGAATTGG CACATGAGAG TGAGTGCCTC CTTAATTTTG AGTGCTGGAC ACTTCTTGCT
225841  CACTTAGCAT ACCCCTGGAC AATGAAGTGT TTTTTGTTTT GTTTTTTCAT GTCCATCCTT
225901  TATCCTTCTT CATCTCAAAA CATTTCAATG GAGTATTTTT TTGGAGCAGT ACTTGGATGA
225961  GCCTCTGAGT CCCACAGTAG CTGAGAATTT ATTTCATAGT ACTCTTTATG ATCACTGTGG
226021  AGCCTTAAAA CATTGTAATA TTAACTTAGC TGGGAACAGA AATTTTGTTC CACAATTTGT
226081  CTTATTCAGA ACAGTATTGA CTTCCTGCTA GTCTCTTCTG ATGTCCAATA TGAGGAAGTC
226141  TAGTTAGCCA GCTACTTTTT GTAGGAGAGC TATGTTTAGG CTAGGTGCTA TAGGATTCTC
226201  TTTATCCTGG AATTCCTTCA CCAAGATGTG CCAAGGTGTT AATCATTTTC TCTTGCTTTT
226261  TGGCTGGTGG TCTTAGAGTT TCCTTCGATT TTGTTTTATT TAGTGATTGT CCTCAATTTG
226321  TTTTCTTTAC TAAGAATCTC TCTTCTATTT ATCTGTATGG TAAAACCTTG TTGCCCATCT
226381  TTCTGGTTTC TGCTGACTTT CATTTTTGGA CCTTTTACTT TGCTTTCTCC ATGGACTTTT
226441  TGGTAGTGGA GGCAGGCAAA CACTTTCCAA AGTCTTTCTC AATTTCCATC AATTTCAACT
226501  TATTTCCTAA AATTGCCTCA GAATGTGCCT ATGTCCACAA TATCCCTCCT TCCACTTTAG
226561  AAAGGAAAGG CATCCACACT TTATTTAGGT GCAATGCCTG AAGTGTAAAC ACTTTCTGGT
226621  TGTCAACAAA GGAGTACTTC CAAATATTGG TTTGGGGATA ACCTGCTAAT GATTAACACA
226681  TTCACCTTGG CTCTTGGTTT GCCTGCTCCC TCTTCTTTTA TCTGCTGTGT GTATTTTTTT
226741  TAATCACTGA GAATATGCAC AGTATTGTAT GTTTTATTAT AAGAGAGGAC TGGCCAGAGT
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226801   GGGAATGTTC TGAATTCAGA ATAACTGAAG CAGTACAGGA TAGGAACTCA TTCTTTCAAA
226861   TGAAGCTGGC ATATTTTCCC AGAGCACCAA ATTTCAATAT ATATTTAAAA AACTTGATAT
226921   GAATGATACA ATAAAGTGGT TAGAACTTTT ATTAAAATAA ACTTATGTCA TGAAATACTT
226981   ATTCTAATTA TAGTCACTCT TCATCTTATT TCATCTTATA ACATGTTTAA TGTTTTCTTT
227041   TATTTACAAA ACAATTTATT TTTTGATGAA AAGTTTTAGA AATCAAGTTA AAAATATTCA
227101   AAGGAATGCC TAAAGTTTTC AAAATTCTTT TACATGTTGT ACAATCAAAA GAGTCTGAAG
227161   ACCATTTAGC TATCCAAATT GTTTATTTTT AAGCAGTATC CCTTCTAATA TTTACTATTT
227221   ATAATCCTTA AAAATTTGCC TTAGCACAGG AGAATTGCTT GAACCCAGGA GACGGAGGTT
227281   GCAGTGAGCC AACACAGTGC CACTGCCCTC CAGCCTCGGC GACAGAGTGA GACTCTGTCT
227341   CAAAAAAAAA AAAAAAAAAA AAAAAAAAG GCCAAAAACA AATAAACAAA CAAAAAAATC
227401   CGCCTTAACA TTATTTGTTC ATTAAAAACT TTCTTTAATA CTACTAGTTT CCCTTTCCTC
227461   TCAGCCCATT GTCATATTTT GATTTTATC ACTTGCTTTG TAGGACATAT GAGGTTTTTG
227521   TTTTTTTTTT TTTTGGAGA TGCAGTCTCC CTCTGTTGCC CGTGCTGGAG TGCAATGGCG
227581   CAATCTTGGC TCACTGCAAC CTCTGCCTCC TGGGTTCAAG CAATTCTCCT GCCTCAGCCT
227641   TCCAAGTAGC TGGGATTACA GGCACCCACT ACCACGCCTG CTAATTTTT GTATTTCTGG
227701   TAGAGACGGG GTTTCACCAT GTTGGCCAGG CTGGTCTCGA ACTCCTGACC TCAAGTGATC
227761   CACAATCCTT GGCCTCCCAA AGTGCTATGA TTACAAGCAT GAGCCACCTG CCCAGCCAGA
227821   ATATATGTTC ATTTTGAGTC CTTTAACAAA GTCATAAGAA TTTTAGGAAT TCAGTTACTT
227881   TCTTGAGAAA ATCTCTGAAA AGATGCCAAT AATTTGTAGC CAATTATATT GATTTCTCTT
227941   TTTCATATTG AGAATTGTTT TTTAAAAAGT TTGTATGTGT GAAGATTTTT GCACTGTAGT
228001   TAAAGAAACC ACCTGTGTGT TGGTTAAGCC ATAAGTACAT GTATTCAAAT AAAATTGAGGT
228061   GGGGTTACTC TGAGAATCAA AGGAAAACCT GAAGAAACAG GCAGCCTCAA AAGGTCTTAG
228121   CTGTAGCAAC TTGCTCCATT GTTGAAATAA ATAGGCTTGA ACTTGTATTT TCCCTCTACT
228181   CAACATTTAA GGTCTCAGAA GATAATATAA TTGGTGAAAT TTAAGTAAAG TGCTCACTCT
228241   TTTGCTTTAA CAAACCCTAG AGAGCTGGTA GGCAGAGCCT CAACAGACCG TTTTAGCTTC
228301   CAAAGGGAGT TCAGGACACC ATGATTCACG ACCACAATAC ATCACACATA ATTGAGAAAA
228361   GATAGTTCCA CCAAATAAAG TTGAAATGCT GACAAGAAGG GGTAAGAAAT CTTGGAAATA
228421   AGTTTATATA AAATTTATTT TTTCCTTTTT TATTGTTATG GAATAGGACC AGTTCTACTT
228481   AAGCCACCCA TTTGCCAAAA TAAAGTGAGA ATCGTTTCTT TTGGGGACTC CTCTTTGTAG
228541   CTCCAAGTGC CACTAACAAT TCTTAGGACC TGAGCTATAA GCCAGGTGAT TTCAGTTAAT
228601   ATGATCAATT ATTTCATTTA AATGGCTCTA ATGTGCAGAG GGAACGGAGC CCATCAGCAT
228661   TCCCTGCAGG GAACTGCAGT GGCTTTTATC AACTTGAACA GCTAGCTTTC AACTGTTTTG
228721   AAATCACTTT CAGGGTGGTC ATGTAGTTGC TTTTTTGAAA TCAGAAGATG ATTCTGCCTC
228781   TTTTAATATG TGACTCCTCA GATTCAGAAA GTGCTCGCTA GTCTTAAGAG TGAATTACCC
228841   TCAGTGGTCC AGCGCTTATG AACCCACATC TAACCCTATC CCCTGGGGGA ACTATCAGAG
228901   AAATTGGTGC CATGGACATA AGAGGAAGGC ACAGTGAAGC AGAGAGCCCC GCATGATGAA
228961   AATCAGTGGA CAGCATCATT ATTTACAACT TTGTAATCAC CCAGGAGCAT GAAAATCCAG
229021   GCCAATCTGG CACCATGAGC TCTAATTTTT GTTGGAGTTC TTGGAACCGA TTCTGATGAA
229081   TGACTGTTTA GCCATTTTAG AGTGTGGCAT ACGTGGCTGC TGGCATACAG AGGTTGGATG
229141   TAAACGGGCC TTTGCCCTCT CTTATGAACA TAGACAGGAA CTAAACTGTG TCACATAGGT
229201   TCCAAATGGT GGCCTGAATA CTATTACAA CTAAGGTACA ATGAAATTGA GTAAGTCTTT
229261   TCCTCTTTTG CAGATACCAT CATTATTCAT ATATTTCTTC AAAGTTAACT ATTTGTATTT
229321   GGTAATTTTT AATAGAAATG TAATAATTGC TTCTCAAGTT TAGTCTTTAG TCTTAAGGTT
229381   GATGCTCTCC ATGTCCTTCC AAAAAAAGGT ATGTTGCTTT TATTATATCC TCGCCTTCAG
229441   ATGGGATTAT TCCATTTTGT TCTTTGTTAA TATATACTTT GAGCCACTTT TTTTGTGGCT
229501   CTGGGTGAGA TGCTATAGGT ACAATGACAA GTGATACGTG TGTTGTCCCT GTCACAAAAG
229561   TGGATAGCCT AAGTGGTGAC TTTTACCTCC ACTCCAAATA TATGTATCAC ACACCAGCCG
229621   TATGCCAGGC ACCACTCTAG GTGCTAGGGA TACAGCAGTA AACAGACAAA TGCAACCCCT
229681   GCCCATGTGA AAGAGAATAA GACAATAAAT AAGTAAAGTG CATGTTATAT GGAGGTGGCA
229741   AATGCTAAAA AGAAAAATTA AGCAGGCAAG AGGACTCATT GAAAAGATGA CATTTGGGTA
229801   AAAGCCCATG TATATATGTT CTATTGGTTT TATTTCTCTG GAGAGCCCTG ACTAATACAC
229861   AATGACTTTG AGAAGTTACT GGCTTTTGAT TTATCACACT ATTCGGAGTG CTGAGAGCCT
229921   TCTTAGTGTG TATTCAGTGT TTTAAGAGAG CTTGTGGATG AATAATAAAT AGGACAAAAT
229981   TTATCCAAAC TTAAGCCTTG CTTTAGGTAA AAGGGCTCCT CTTACAAGGT AGAAGGTTAT
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230041    TATTTGGCAT TTAAATCCAA CTGAAGACTA ATAAGACTAA TTAATTAAAA GTTTTTAAAT
230101    CACAACTGGG TGCAAAATAA ATGGAACTGC CATGCTCGCC AAGTGTGCAT GAGTGGTGTG
230161    CATGGGAGAC AGCACGAAGC TAATCCCACT CATCTTGCAG GTTGCTCCAT TTTTCTCCTA
230221    AAATCAGTAA GACAGAAGCT GGTCAGATTA TCAAGAGCCC TAGTTAAACA CAGCAGTAGC
230281    ATTTGGAAGG GGTTGCTCTC ATTAGGCAGT GCCTGACCAC AACAAGAGAT GAACAAGCCC
230341    TGTATCTGAA GCCATCATGC CTAGTTATGG TCCCCCACTG TTCATGATGC CTGAAAGGGA
230401    GGCCCCCTGC ACCCTAGAAA GCTGGGTGGG TTCTACTGTC TGCTTTACTG CTAAAAACCC
230461    TCTTCTTTGG ATCTGGACTT TACCTCTATC TGATTTTTTT TTCTAATATA TGATTTGGCA
230521    CTGAGTCTGT CACTGCTGCT AACTCAGCAG TTCTAGGGTC ATTGCCCCAT TGCCTCACAG
230581    AAAGAATTTC ATAGCTTCCA GCATCCTCTC TCCTTCATTA TACTTTGATT TCAGCATTGC
230641    TATTTTTTCT CTTGGGTGTT GCAGCTCTCT CTCTCCTTCC CATGTCTTGT TGGTTTTCTG
230701    CTAACTCCTG CTTTTTTTCT TTTTTTTTTT TTGAGACGGA GTCTCGTTCT GTCACCCAGG
230761    CTGGAGTGCA GTGGCACAAT CTCGGCTCAC TGCAACCTCC GCCTCCCGGG TTCAAGCTAT
230821    TCTCCTGCCT CAGCCTCCCA AGTAGCTGGG ACTACAGGCG CTCACCACTA TGCCCCACTA
230881    ATTTTTGTAT TTTTAGTATT GCTGTCATCA ATCCACATGT CCAGAAGCAC CTAGAAACTC
230941    TAATTCTTTG TAGGTATCAA ACCCTAGGAC TCTTTCCTCT AATCACAATA TATAATCCCT
231001    GATTCCCAAA CACGGTCTTT TCATATACAT TTTCCACTGT ACATACTTTC TGACCTGGAA
231061    AGCTCTTACA CAAACACGCC CTCCCCTAGG AAGCCTTTAT AAATGTTCCC AGGAAGAATC
231121    AGTCACCCAA CAGTGTCCTT GTCACATCTT AGGTTCTACA CCTTTATTTG TTCTATCTGA
231181    ATGTAATCTC CCAGAGGGTG TTATCATCTT TTTTTTTGAG ATGGAATCTT GCTTTGCTGC
231241    CCAGGCTGGA GTGCAGTGGC ATGATCTCGG CTCACAGCAA CCTCCACCTC CTGGGTTCAA
231301    GTGATTCTCC TGCCTCAGCC TCCTGAGTAG CTGGGATTAC AGACGTGTGT CACCACACCT
231361    GGCTAATTTT TGTATTTTTA GTAGAGACAG GGTTTCACCG TGTTGGCAAG GCTTTCCTCG
231421    AACTCCCAAA CTCAGGTGAT CCACCCGCCT CAGCCTCCCA AAGTGCTGGG ATTACAGGTG
231481    TGAGCCACCA TGTCCAGCCC CATCTTTTTC TTTTAGTTTA GTTCTTAACA AATAGTCTGA
231541    CACAAAGTGG ATATAACAAT ATTTTGAATT ATGAATAACT AAATGAATAT TTCCAGATTT
231601    CCTGGTGCTC TCAAAGTTTT ATGTTACAAA AGAAAAACAA GTCTAAAATA CCTGCCTCAA
231661    GTTTTTATCT GTACTATGAT TTCAAACCAA ATAAAAAACA GGTGGGGTAA AAACTGAAAC
231721    AGGAAATACA TATAACTGAA AAATTTTGGT ATGTTAGTAT GATAATACTA GGTCATTTTT
231781    CCTGTTTCCC CAACTTCATT TTCTATAGCA ATAAAAAGAA ACAAGTAAAT GTATATTAAT
231841    TTAATTTAAA AGAAGTAGTC TACCATCTCT TCTGTTAAAA AGAAAAAAGT ATTTTAAAAA
231901    ATTATCTCTG GAAGGATACA CAGGGAACAT TGCTCTGGTT TCTTCCAAGA GAGAAATGAG
231961    GAACTAGAGA GCATGGCCAA GTGGGGTTTT GCTTTTGTTT TTGTTTGTCT ATCTGTTAGC
232021    TTTTTATTAT TTTCTTTTGT AGGTTTGAAT TTCAAACCAC ATAAATCTGT TACATGCTCA
232081    TAATAATAAG TTTAAAATAA AACTTTTGGC TGGGTGCAAT GACTTACACC TGTAATCCCA
232141    GCGCTTTGGG AAGCAGAGGT GGGAGGATAC TTGAGGCCAG GAATTTGAGA TCAGCCTGGG
232201    CAACATAGTG AGACCCTGCC TCTGTAGAAA TAAACAAAAA TTAGCTGGAT ATGGTGGTGC
232261    ATGCTTGTAC TCCTAGCTAC TTGGGAGGTT GAGGCAGGAG GATCCTTTGA GTCCAGGAGT
232321    TTGAGGCTGC AGTGAGCTAT AATCACCCAC TGCACTATAG CATGGGCAAT AAGGTGAGAA
232381    CTTGTCTCAA AAAAAAAAAA AGGGGGGGGG AAACAAATAA ATAAATATAA ACAAAACTTT
232441    TGTTCAAAA TATGTAATAT TTAGCACTAA AGAATTCTGA ATTGTAGAGC TAAAAAGTAC
232501    TTAAAGTTA ATAATTATTG TCTCCTTTAA AAGAATTGTT ATCAAAGTAT AATTTTTATC
232561    CAGAAAATCA TCCATATCAG CAAGCTAAAC TTTCTCAAAA TGACATATCC ATGTAATTAG
232621    CTCCCAGGTA ATTAGCAGGC AGCCTCTACT CAGGTTGAGT ATTCCTAATC TAAAAATTGG
232681    AAATTCAAAA TGCTCCAAAA TCGGCAACTT TTTGAATGCT AACATGATTC TCAAAGGAGT
232741    GCTCATGGAA TATTTCAGAT TTTGGATTTT TGGATTTGAG ATACTCAGTA TAATGCAAAC
232801    ATTCCAAATC TGAAAAAATC TGAAATACTT CTGGTTCTAA GCATAAGGGA TACTCAACGT
232861    GTGTTAGCTA ATTAGACCCT TCATGGTCTC TTCTAGACCT CAGCTTCTTC AAGGTAACCT
232921    CTATCCTCAC TTCTAATAGC ATGAACTTTT CTGTTTTAGA ATAATTTGGA TTTTCAGGAA
232981    AGTTGCAAAG ATAGTACAAA GACAGTACAG GAGAGTTCCC ATATATCTTT CACCTAGCTT
233041    TCCCCCATTG TTAGGATTTT ACATTATTAT GATACATTTG TCAAATATAA GCAACTCACA
233101    TTGATACATG AAACTCTATT AACCAAACCC TAGACTTTAT GTGGATTTCA CCACTGTTTC
233161    CACTAATGTT TTCTTTCTGT TCCAAGGTCC AATCTGGAAT ACCACACTGC ATTTTCTTGT
233221    CATATCTCCC TAGTCTTTTT TTGTCTGTGA CAATGTCTCA GTCTTTTCTT GCTTTTCATG
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233281   ACCTTAACAG TCCTGAAGAT CATTTGCTTT TTTTTCATAA TTACACCGGA GTTATAGATT
233341   TTTTGAAATA ATACCACAAG GGCAAAGGGC CCTTCTTGTC ACATCATTTT AGGGAGAACA
233401   TGATATCCAC ATGACATCAC TGATATTAAC CTTCATCATG TGGTTTAGGT AATGTTTCAG
233461   GTTTCTCTAC TGCAAAGTGA TTTTTTTCCC TTAATTTAGC CCACCTGAAC TTATCAATTT
233521   TGTTTTCTTC CATGACTAAT ACTTTTGTTA TTATAGCTAA AACTTCATTG GGGCCAAATC
233581   TTAGATCATG TAAATTTTCT TCTATATTTT ATTCTAAAAG CTTGTAATGT TTGATACATT
233641   CTAAAAGATG TAATGTTTGA TACATTACAT CTAGTCCTTT GATTTATTTT TAGTTACTTT
233701   TGTATAAGGT GTGAGAGATG TCTCCAGTTT CACTTTATTA ACACATTGTG GTGTTCCAGT
233761   ACTATTTGTT GCTAAGACTA TCTTTTTTCC ATTGATTACC TTTGCCTTAG TTGGCAATAT
233821   TTTTGTTGGT TTATTTCTAG ACTGTTTATC TCATTCCACT GATTTGTGTC TATCTTTTTG
233881   ACAAAACTGT TGATTACAGT AAGCTTTGAA ATAGTTCATT TTTTGTGTCA ACTTGACTGA
233941   GTCAGGGGAT AACCAGCTAT CTGGTTAAAC ATTATTTCTG GCTGTGTTTG TGAGCGTGTT
234001   TCTGGATGAG ATTAGCCTTT GAATAGGTGA TCCTAGTAAA GTAAACTGTC TTTCCCAGTG
234061   TGGATGGCAT TATGCCACCT GATATTCAGG GTCTGAATAG AAGAAAAGGC AGAGGAAGGG
234121   GGAATTTGGG CCTTTTTTTC TGCCTCACTG CTTGAGCTGG GACATCTCAT CTGGTCTCCT
234181   GCTCTTGAAC TGGGATTTAC ATCATCAGTT CCTCTGGTTC TCAGGCCTTC AGATTCAGAC
234241   TGAATCATAC CACCAGCTTT CCTGGGTCTC CAGCTTGCAG ATTACAGATC ATGGGACTCC
234301   TCATCTTCCA TAAATGCATG AGCCAATTCA GTCTATGTCC TTGAAAACTG CCCCACTGCA
234361   GATTAAGGCT TTTTTCCACT AGGTGAAATA AAGAAGCTTG TTAGACAGAT TTCCCTTCAT
234421   CCAGTGCCCT CTCCTCTTTA AGTTACAACA CATTGGCTAC ACCTAAGTGC AGGGGTGGGG
234481   ATGAGGGTAT AGTCCTCTTG TTTGCTGAGA AGAGAACTGT ATTGGGAAAG CTCTAGAAGT
234541   GTTTGATACA TACATAAACA AGGCATGGTT TTTGCACTTA ATTTCACATT ACATTTTTCC
234601   CAGAAAAAAA GGAATGTATA GGCATCACGT AACTGTACTA GCTGGAGTCA TTCTTCCTGA
234661   TTATCAAAGG TAAACAGTTA TTAATCCTAT ACCAAGATGT CAAGGAGAAG TACTTTTGGA
234721   ACACAAGGAA TTCTCTGGGA GTCCTTACTA CTCTCAAGCC CAGTGAAAAA GTTAATGAAA
234781   AACTATAGTA CCTTCCTATA AGCTGGATGA CTAATTACCA GGCTCATTTA GGAATTTGCC
234841   TTACCAAGTA AAACATAAGG GCAGCTGAGG TGCTGACTGA AGACAAATGG AGCATAGAAT
234901   AAGAGTAGTA AAGAATGCCA AAAATGCTGT CATGTATCCA TTGACAAAAG GAGCTATAAA
234961   GCCTTTAGGT ATTTTCACAC TTGCTCTGTT ACGTAAATGT ATGTGTGTGT GTGTGTGTGT
235021   GTGTGTGTGT GTG
```

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   1   CACACACACA CACACACACA CACACACACA CACAAATGAG GTATATAAAG GGTCTCCTAA
  61   AATGTCATCT GATATTGTT ATTTCATATT CTCAGATTTT TAATCCATTT AGGTAGGTCT
 121   ATTTTAGATA GCCTTGTCTG AAACAGAGCT GGGACCTGAT GAGTGAAAAT GAGCTCACCA
 181   GAAGAAAAAT CAAACAGGCA TTTCAGAGAT TGAGGCCAAG AAGTTAAATG TCTTAAATGG
 241   GCAGAGCTTA GCTGCTTGAT GTGAAAAGAG ACCAGCGTGG CTGGAACAGC AAAGGAGAAC
 301   AGCAGAAGAG GTGAACAGAG GCCAGAGATG GTCACTGAGT GGGCCCTTAA GTCATGGTAA
 361   GGAGTATGGA GAATGAATTA TTGCATGTAT TGAATATGTA GGTGACGTGA CTCACAGATA
 421   CTTTGGATTT GTAGAGATGA AGGAAATGTA GCAAGTGACA CTCTTAGAAT GTTGATTTGA
 481   GTAAATGGTA GTGTCAGTTA TTGAACTGGG GAGAACTGGA AGGGATAACA GGCTTAAGGA
 541   GCACGTTTAT TCCTGTGTCT TGGAAGTGTT TAGGGTGAAA GACCTATTAG AGTTCTAAAT
 601   GGAGATGTCA AGTGAAAATG TGGCTACACA CATTTGCATT TCAGAAAAAA GGTCAGGCTG
 661   GAGATGTAAA ATTGGAAGTT TACTGCATAT AGATAGTCTT TGGAACCGTA GTATTGATGA
 721   AGCCATTAAT GAGACAGAAC AAAGACTAGG GACCAGAGCC AAGCTCCAAG TTTCTAAAAT
 781   TTAGAGGATA GTATAGTCTG GTCATTTTGA GGTGAATACT TAATAACAGA ACAATTTGCT
 841   GAAGTGTAAA TTTAGAGCCC TACACTTTTA GCTCTGACTA TTAACGAATA CAGGAAAGAA
 901   TGGATATGGT TATCTGCCTG GTGTCTGTGA AATAATTTAA GCCAGGAAGA GATCCTCACC
 961   AGAAACTGAC TATGCTGGCA ACTTGGATCT TAGATTTCCA GCCTGCAGAA TTGTTAGAAA
1021   ATAAATGTCT ATCGTTTAAG CCACCAGTCT GTAGTATTTT GTTATGGCAG TCCAAGCTGA
1081   CTAAGTTTTG GTACCCAGGC GTGGGATGCT GCAACAACAA ATACCTAAAC ATGGGGAAGT
1141   GGCTTTGGAA ATTGGTGATG GGTAAAGGCT GGAAGAGTTT GAGGTTCATA CTAGAAAAAG
1201   CCAATTGTGA AGGGACTATT GAAAGAAATA TGGACATTAA AGGCAATTCT GGCAAAGGCT
1261   CAGAAAGGAA GAGAGCTGGA CAGAAAGCTT CCATTTTCAT AGAAACTTAG ATTTATAACG
1321   ATCATGGATA GAATATTAAA TATGCTGGTT AAAAATATGGA CTTTAGGCCA GGCGTGGTGG
1381   CTCACGCCTG TAATCTCAGC ACTTTGGGAG GCTGAGGGCA CAGATCACGA GGTCGGGAGT
1441   TTGAGACCAG CCTGGCCAAT ATGGCGAAAC CCTGTCTCTA CTAAAAATAC AAAAATTAGC
1501   TGGGCATGGT GATGTGCTTC TGTGGTCCCA GCTACTCGGG AGGCTGAGGC TGAAGAATCG
1561   CTTAAACCCG GGGGGTGGAG GTTGCAGTGA CCCAAGATCA CACCACTGCA CTCCAGCCTG
1621   GGATACAGAG CAGGACTCCA CTCCCCCCGC CACACACACA CAAAAAATAT ATATATATGG
1681   ACATTAAAGT CAACTCTTGT GAGGTCTCAG ATGAAAATGA GGGACAGGTT ATTGGAAACT
1741   GTAGAAATCA CTGTTCTTGT TACAATGTGT CAAGAACTTG GCTGAATTAC GCTGTAGTGT
1801   TTACTGGAAA GAACTTATAA GCAGTAAAAC TGGATATTTA CCAGAAGAGA TGTCTAAGCA
1861   AAGTATTGAA GGTGTGATTT AGGTCCTCCT TACTGCTTAA AGTGAAATGT GAGAGGAAAG
1921   AGCCGAAATA AAGAAGGAAT TTTTAAGCAA AACACAATCA GAACTTGGAG ATTTGGGATA
1981   GATTTCTCAA TCTATATTGT AAAAATTGAG AAAGTTTTTC TTGAAGAGGT ATGGTTGAAC
2041   AATGTTTTCT TTTTCTTTTT TTTTCTTGGT TTTATTTTTA TTTTTATGTT TTTTGAGACA
2101   GGGTCTGGCT ATGTCATCCA GGCTGGAGTG CAGTGGCACA ATCTCAGTTC AGTGCAACCT
2161   TTGCCTTCAG GCTCAAGCAA TCCTCCCACC TCAGCCTCCT AAGTAGCTGG GACTACATGT
2221   ATGCACCACC ACACCCTGGC TAATTTTTTG TTGTTGTTTA TAGAGATGGG GTTTTGACAT
2281   GTTGCCTAGG CTGGTCTCTA ACTCCTGAGC TCAAGTGATC TGCCCTCCTC AGTCTCCCAA
2341   AGTGTTGGGA TTACAGGCGT GAAACACTGA GCCTAGCCTG AACAACCATT TGATAAAGAG
2401   ATAATGGGTG TGACCCAAGG ATTTAATCAG CCATCTCAGC AGAAGCCAGG AAGAGAGATG
2461   GGATTATTCC AGCAGAGACA CTGCCAATTT AAACTAACGT AGGCAGAGAA AACAGAAAGG
2521   AACAAAGGAA GGTTGTCGAC TTTTTGAATT CTATAGAACA GGATCATAGA GCTACCTGGC
2581   TGTCAATGTG TACTATTCTT TAAGAAAAGG AAAGACTGAC CCACCAAAGG CAACTTACAA
2641   GATCACTAGG GCTGACTCTT TTTTGTTTTT TCTTGAGGCA GTCTCACTGT CACCCAGGCT
2701   GTAGGGCAAT GGTGTGATCT CAGCTCACTG CAATCTCCAC CTCCCAGGTT CAAGGGATTC
2761   TCTTGCCTTA GACTCCCAAG TAGCTGGGAT TACAGGCTCT AAATCTGTAC CCTCCCGAGT
2821   AGCGCTCCTG CCACCACTTG CCCAGCTAAT TTTTGTATTT TTAGTAGAGA TGGGGTTTCA
2881   CTATGTTGGC CAGGCTAGTT TGGAACTCCT GACCTCCAGT GATCCATTCT CATTGGCCTC
2941   CCAAAGTGCT GGGATTACAG GCAGGAGCCG CCAGGGCTGC CACTTTGATG TCAGACTCAG
3001   AGAGTACAGA TGGGATAGGG TGGGGGTGGG AACATGTAGT CAAGGCTGAC TCTACCTGTT
3061   TCAAAGATGC CCTGCAGAAC TGTGTGGGAG TCTCTCACAG ATGGCTGCCT GGGTGGGACC
3121   CCACCAAACT GAAAGACCGA GACTTCAGGC AGGGCAGATG GAGTAGGCCA ACTACAGAGC
3181   CAGAGGTGAC ACTGAGACAC CACTGGGCCT GGAAATCAGG GCATCAAGCC AAAGAGGGTT
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3241  TTTCTTAAGA  CCTAACAGAA  TTTGCCTTGC  CAGGTTTTGG  ACTTGATTAG  GACACATTAC
3301  ACCTTCCTTC  TTTCCTATTT  CTCCATTTTC  TAATGGGAAT  GTCTATTATG  CCTGTTTCAC
3361  CATTGTACCT  TAGAAGCATG  TAACATTTCT  GGTTTCACAC  GTTCAAAGCT  GGAAAGGAAT
3421  TTTGTCTCTG  GATGAATCAC  ACATTGAGCC  TCACCCGTAA  CCTGATTTAG  ATGATTTTTT
3481  AGATGACACT  TTGAACTTTA  GAATTGATGC  TAGAATGAGT  TAAGACTTTC  AGGGGGCTGT
3541  TGGGATGGAA  TAATTTTTTT  TTTTTTTTTG  AGACGGAGTC  TAGCTCTGTC  GCCCAGGCTG
3601  GAGTGCAGTG  GCACCATCTT  GGCTCACTGC  AAGCTCTGCC  TCCCGGGTTT  ATGCCATTCT
3661  CATGTCTCAG  CCTCCAGAGT  AGCTGGGACT  ACAGGCGCCC  GCCACCACGC  CTGGCTAATT
3721  TTTTTTTTAT  TTTAGTAGAG  ATGGGGTTTC  ACCGTGTTAG  CCAGAACGGT  CTCGATCTCT
3781  TGACCTTCTG  ATCCGCCTGC  CTTGGCTTCC  CAAAGTGCTG  GGATTACAGG  TGTGAGCCAC
3841  CATGCCCGGC  TGGGATGGAA  TAAATTTATC  TTGTATGGGA  GAAGGACATA  CATTTTGGCA
3901  GGTCAAGGAC  AGAATGTTAT  GGACTAAACT  GTGTCCCCCA  AAATTCATTT  ATTAAAACCC
3961  TAAACCCCAG  TGTGACTGCA  TTTGGACATA  GAGCCTTTAG  GGGGTACATA  AAACTAAAGA
4021  TCACAGGATA  GGGCCCTAAT  CCCATTGGGG  CTGGTGTCCT  TACAGAAGAT  GAGACACTTA
4081  GAGCTCTCTC  TCCACGCAGG  CACCAAGGAA  ACACCATACA  AACACACAGT  GAGATGGCAG
4141  CCATCTGTTA  GCCAGGAACA  GATTCTCACC  ATAAACTATG  TTGGCACCTT  GATCTTAAAC
4201  TTCCAGGCTC  CAAAACTGTG  AGAAAATGAA  TTTCTGTTCC  AAGCCTCTTA  GATATGGAAA
4261  AAAAGATTCT  GTTGTTTAAG  CCATCCAGTC  TCTGGTATTT  TGTTATGGCA  GCCTGAGTAG
4321  GCTAAGACAA  TGAAGGATGT  GGTAAAACTT  TACGTCCCAA  CCACATACCA  AAGAGGCTGG
4381  AATTTAGCAT  GCTTTCTTCT  TTCAACTGTA  GGCAATGTGC  ACAAGTTCTA  AATCCTAAGA
4441  CATGTTGGCT  CCTTTACTCT  GCCCAAACTA  CAACTCAAAC  AAACAACTGT  AATATAATAA
4501  CATCCAATGA  AGTTCTGACA  TTTCTTCAAC  ATGAGTACAG  TAATTCAATG  CCAGAGAATT
4561  CATTTTATTT  TGAAATCTAC  ATGCCATATT  CCAATTTCTG  TTGAAGATGC  AATGGTTATA
4621  TTTATTCTTT  TTAATATAGA  TTTATCAGAC  TGGGCGCGGT  GGCTCATACC  TGTAATCCTA
4681  GCATTTGAGA  GGCTGAGGTG  GGCATATCAC  CTGAGGTCAG  GAGTTTGAGA  CCAGGCTGGC
4741  CAACATGGTG  AAACCCTGTC  TCTACTATAA  ATATAAAAAT  TAGCTGGGTG  TGGTGGTGCA
4801  TGCCTGTAGT  CCCAGTTACT  AGGGAGGCTG  AGGTAGAATT  GCTTGAACCT  GGGAGCAGGA
4861  GGTTGCAATG  AGTGGAAATC  GCACCAGTAC  ACTCCAGCCT  GGATGACAGA  GCAAAATAAT
4921  AAATACATAA  AATAGATTTA  TCAGTTTATC  AATAATATAG  TTTTCTTTTC  TAGGTGTAAA
4981  TATAGGTAAT  GACTGTCCTT  TAGTACATTT  TCTCATGATG  CTCCTCTTAC  TTGGTTTGGT
5041  ACAATATTAA  GTATTGAAAT  AAAATAGAGA  ATCCTGTCGC  TACACATGAG  CACTTATTCC
5101  ATTTGCTCAT  CTCCAATATG  CACGGGAAAT  TCTCAAATTG  CTAATAATCT  TGTAACACAC
5161  ATGCATTATA  TTCAACAGGA  ATATATAAAT  TTATAATTAT  AATTTAGGAT  CAACAGATGA
5221  CAAACCTTTA  GAAGGTTTGT  ATTTAACCTT  AAAATATAAT  TTTTTAAAAA  TTGGTTATAA
5281  AATTTCTAAT  ACTTTCTTTT  TTGTGACCTC  AAGGGGAAAA  TATAATTCTT  ATAAAAGTTC
5341  AAATGATTTA  CAGAATACAA  AAAGTGAATA  GAGATGATGA  ATGAATTAAA  GGAAAGGATA
5401  TTGCTACATA  GATTGGAAA   TTTAAAAAGG  GAAATTACGA  TTGTTGATTT  TGTGTTAAAC
5461  TGATCTGCTT  TGTTCAAGAT  ACCTTATGTA  CCAAAAAATG  ATTTTATCTC  AGCCTCATAT
5521  CTCAGTAAAT  TCCTGAGACA  AACTTTAGTC  CCTGGTGCCC  AGGTGCCTTT  GGTAATTGGG
5581  AGACCTCTAG  GTTTAGCATC  CTCATCCACT  CGCCCCAATT  TAAATAGTCC  TCCCCAGGGC
5641  CATTCAGGCA  AGGGAGATGA  AAACTTGCTC  AAGAGTTGGA  ATCCAATTGA  AGCTACCGAA
5701  ATTCATTGCT  CAATAGATAA  TTTTCCCTGG  AAGTAACTAG  GGCTTTTGAA  TATAATAGTG
5761  GGCATTTCAA  AGTAGAAGGT  AAAGTATTTT  GGAGATGAGG  AGACAGGACA  GAGCTACGAG
5821  GAATGTCCTT  TGCTCAGGGA  CTAGGCTCTT  AGCAGTACCT  CTTAGGTAAG  AACTGGTTAA
5881  CTGGCACCTT  CTGTGTTTCT  CTGAAGCTCC  CTTTGCTTAG  GGACTAGGCT  CTTAGCAGTA
5941  CCTCTTAGGT  AAGAACTGGT  TAACTGACAC  CTTCTATGTG  TCTGAAGCTC  CCAGAACAAA
6001  CTGCCAATGA  AATTTGGATT  TTTGGAATAT  AGTTTCTTTT  TTGTTGTTAC  TTTTTGTTTT
6061  GTTGTTTTTT  TTTGAGAGTC  TCACTCTCAC  TGCAACCTCC  CCTCCTATA   TTCAAGTGAT
6121  TCTCTTGCCT  CAGCCTCCCG  AGTAGCTGGG  ACTACAGGCG  TGCACTAGCA  TGCCCAGCTA
6181  ATTTTTGTAT  TTTTTAGTAG  AGATGGGGTT  GGTTTTTTTT  TGAGACAGAG  TTTCACTTTG
6241  TCGCCCAGGC  TGGAGTGCAG  TGGCACGATC  TTGGCTCACT  ACAACCTCCA  CCTCCCGGGG
6301  TTCAAGTGAT  TCTTCTGCCT  CAGTCTCCTG  AGTAGCTGGG  ACTACAGGCG  CCTACAGGTG
6361  AACACCGCCA  CACCTGACTA  ATTTGTGTAG  TTTTATTAGA  GATGGGGTTT  CGCCATGTTG
6421  GCCAGGCTGG  TCTCAAACTC  CTGACCTCAG  GTGATCTACC  CACCTCAGCC  TCCCCAAGTG
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6481   CTGGGATTAC AGATGTGAGA CACCAGATCA GCCTCAGAAG ACATTTTCTA TTGGAAAGAG
6541   AAAACACTAT TAGCAACCTA TTAGTCTAAT ATTTAATACT TAATGTCTTC CTTAGTAATA
6601   AACCAACTCT CTACAACAAA GTGCTTCCTG GCTGCCTAGT CATTGATTCA TTCAGTTCAA
6661   CATTTTCTCA ATGCCCAACA GCCAAGTGTC TCCTGTATGC CAAGTTCTAT GCTGATTATC
6721   AGTATTTGAA TAAGAGGGGG TCTACATCTT AAGTACTGCT TAAGATGAAA GCCTCTAGGT
6781   TAACAAACTT AACACAATGT ATCATTCACT ACTAAATAGA CCGAATACAA AATCTTGTTA
6841   TTGGAGCCCA GAGAGAAGAA TTGAAATTCA AGTTTTCTCT CTCTCCTTTT CTCACTCACC
6901   ACAATAAGTC AGTTGCACCA AGTCTTGTAG CTCTTTACTG AGCCATGTTT TCACGTGTCC
6961   CTTTGTTTTA TTTGCCACAC CCTAAATAAA AATTGTACTG GCTTTTTTTC CTGGGTTTA
7021   CAGTATTAAT ACATTGTCAA GATTTACCTC TTCGTGTAGA TTCCCTGGGG AAAATTACCT
7081   TTCCTCCTTC CCTTAAATTC TTCAGAGGTT AGAAAGCCAT TAGTAACATT CTGGTATGTG
7141   GACAAAGTTT ACCCATTATG TATGGATGTT TTACTCTTTC CATTTTTCTG ACAATAATCT
7201   CTTAAGGAGG TGTGGTTATA GAATAGTCAG CTGTTATAAG TACTGTTTTC CTGGCCTTAC
7261   AACTTAAATT CTTTAAGCTG TTTCTTAGTT TGCTCATCTC AAAATTCGGA ATAAGGATAA
7321   AACCTATCTC TTAGATTGTT GGATTAAATG AATTAACATA CTGGAAGCTC ATGAAATGTG
7381   CCTGGCACAC AGTAGTGCCT AATAAACCAT CTCTCTTATT CAGCCTGTTT TCTGATTTCA
7441   GAATCTACAC TTGCTGAGCC AGGTTCTTTT CATTTCAAGG TGAGCAAAAG CATACAAGGA
7501   AGAGATGGAG GTAGGAAGAG ATTAAGCCCT AGGCCAAGGG AGCTGGAATC AAAGGCAATT
7561   TGGTCAGTGA ATAAAAAGGA TTCCAAGGCC CATAAGGCAA TTCTAACCTT AGGATCGAAA
7621   TTCTCGGACA TACAGGAAAT GCTGGGGGGG GGAAAATCCG GTCTTCTCAG CCCAAGAGCC
7681   ATGTGAAACC AGACCTTCAA ATCTGATGAT TCTCAGCCCA GCTGCCCATT AGAATCGTTG
7741   TAATTTAAAA ATACCCTCGG AAAATTCTAA TATGTGGCTA TCAAAGGTGA TCATTTGCTT
7801   TTATGCCACT TTGTTTTCAC CCAAATGGGA CATCCAACCC TTTTCCTTTG AGAGTAGTTG
7861   TAGGGAAAGG AGGGGGTGGA GGGAGGGAAG AGCGGAAAAG CTGGATCCG CCCCGAGCCG
7921   GTGTCAGTAT CTGGGAAGTG GGAGGCGCGT CAGCAGTAAA CAGCTTCTGC TAGGATTATT
7981   ATCTCCTGCC ACACACTCGG ATTTGAAGGC TCCAAACGAA ACAATGCAAA ACGCTTCAGT
8041   GGAGTTCCAG AAGCGTTAGA CTAAACGACT GGGTCTGTTT GGCCAGTCTG AGCAGCTGGG
8101   CGCAGATGCA TAGGCAAGAC TTAGCCCGCC TAGACTTTTC TGCCCACTTA ATTCCGATCA
8161   AAGCAGAAAC CGGCCGGGCG CGGTGGCTCA CGCCTGTAAT CCCAGCACTT TGGTAGGCAG
8221   AGGCTGGCGG ATCACCTGAG GTCAGGAGTT CGAGACCAGC CCGGCTAACC TGGTGAAACT
8281   CCGTTTCTAC TGGTGGCGGG CGCTTGTAAT CCCATCTACT AGGGAGGCTG AGGCCGGAGA
8341   GTCGTCTGAA CCCGGGAGGC GGAGTTTGTA TGCAGTGAGC CGAGATCGCG CCACTGCATT
8401   CCAGCTTGGG CAACAGGAGC AAAACTCCGT TTCAAAAAG CAAGCAAACA AACAAAAAAA
8461   TGCAGAAACC GAGATCCGGA AGAAAACCTC GGCGAGATTC ACAGAATCCA GGAAAATAGG
8521   TCTCTAGAAA TTTGTCCATG GTCCCAGATC TCCATTTCTT GTGGGTGGGG CAGCTGTTAC
8581   CAGATCCCTA GAAGCAAAGG TTTTTTTGGG GGACCGTGTC TCACTGTTGC CCAGGCTGGA
8641   GGGCAGTGGC ACGATCTCGG CTTACTACAA CCTCCGCCTC CCAGGCTCAA GCGACTCTCC
8701   TGCGTCAGCT TCAAGAGTAG CTGGGAGTAC AAGGTATGTG CCACCACGCC CAACTTATTT
8761   TTTTATTTAT TATTTTTATT TAGTAGAGAG GTGTTTCACC ATGTTGGCCA GGTTAGTGTC
8821   GAAGTCGTGA CCTCAGGTGA TCAGCCCCCT CGGCCTCCCA AAGTGGTAGG ATTAGAGGGG
8881   TGAGCAGAAA GCAAAGGTTT TTGAGTGGCC ACAGGCCCCA CTCTATTTCC TTTTCTGCCT
8941   GTAATGGCAA CCTAGACGCT TGAGCTTCTT AAAATACAAG AGTAAGTTGC ATGTCAGGCA
9001   CCGTTCTACA TTAGGGACAT TAGTCTGTTT TACAGACACC TTTCAACTCC CTGGTTAACT
9061   TTTAGGTAAT ATACTCTGCA CTTTAGCAGG AATGGAACCT ATAACTCTCA CAGAATTAGG
9121   AAAGTGAGGC TGCCTACAGC CTAAATTGAG AAAAAAATAG ACGGGGACT AGTCGGAGGA
9181   CCAAACAAGG TTACCAACAC GTTAGAGTTT TGCCTTCAAT TTACATTTTT AAAGTAATCA
9241   CAACGAAGTG TTTAGATCAC GAGGCATCCC TGCATGTAAA CTGTTAGGCA CTAACTATGG
9301   TCGATCTTAC AAAGCATTAA CTAGAATATT TCTTTAGAGT ATGATAGTAC GTAACTGACC
9361   TACTATTACA TACAAACAGA CCAACCTTTA GTAACAGCGC TCCCCAAAAA CCGAAAGCA
9421   GTAATACGCT TTGCTCAAGG TTGGCATAAA ATTAACTTAC CTTAGTGCCT TTTTTCCTTC
9481   TACCTACAAG CAGTGAGGTT AGCTCTTCCT TTGAAACGGT AGGGGGGCTC TGAAAAGAGC
9541   CTTTGGGTTT GATAGCGTTT CCGGGAGCTC AGATACCTGT CAAATCACTT GCCCTTGGCC
9601   TTGTGGTGAC TCTCGGTCTT CTTAGGCAGA AGCACGGCCT GGATGTTAGG AAGGACGCCG
9661   CCCTGAGCAA TGGTCACCCG GCCTAGCAGT TTGTTGAGCT CCTCGTCGTT GCGGATGGCC
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 9721    AGCTGCAAGT GGCGCGGGAT GATGCGAGTC TTCTTGTTGT CGCGAGCCGC GTTGCCGGCC
 9781    AGCTCCAGGA TCTCGGCGGT CAGATACTCT AACACCGCCG CCAGGTACAC CGGCGCGCCT
 9841    GCCCCAACCC GCTCTGCGTA GTTGCCTTTA CGGAGCAGGC GGTGCACTCG GCCCACCGGG
 9901    AACTGGAGAC CAGCGCGAGA AGAGCGGGAT TTCGCTTTGG CGCGAGCTTT GCCTCCTTGC
 9961    TTACCACGTC CAGACATTGC AATCAGACAA AAATCACCAA AACCAGCAGC CTAAGCTCAC
10021    GAGAAAACAA ACAAAATCAA GAAATATGTA AAACATGGCC GCTTTTATAG GTAGTTCCTG
10081    GGGAGTAAAT CCGACTTTTT GATTGGTCGG TAGCAAATGC TAGTCAGATA GCCAATAGAA
10141    AAGCTGTACT TTCATACCTC ATTTGCATAG CTCTGCCCAC GGATGACAAC TGTGTAGTTT
10201    GTCTTCCAAT TAACTAAGAG GTACTCTCCA TCCCTCATTA GCATAAAAGC CCTATAAGTA
10261    GCAGAAATCC GCTCTTTACT TTCGACACAT TTCTGGTGTT TTAAGATGCC TGAGCCAGCC
10321    AAGTCTGCTC CCGCCCCGAA GAAGGGCTCC AAGAAGGCAG TGACCAAAGC GCAGAAGAAA
10381    GATGGCAAGA AGCGCAAGCG CAGCCGCAAG GAGAGTTACT CTGTGTACGT GTACAAGGTG
10441    CTGAAACAGG TCCATCCCGA CACTGGCATC TCTTCCAAGG CCATGGGCAT CATGAATTCT
10501    TTCGTTAACG ACATATTTGA GCGCATCGCG GGCGAGGCTT CCCGCCTGGC GCATTACAAC
10561    AAGCGCTCGA CCATCACCTC CAGGGAGATC CAGACGGCCG TGCGCCTGCT GCTTCCCGGA
10621    GAGCTGGCCA AGCACGCCGT GTCGGAGGGC ACCAAGGCCG TCACCAAGTA CACCAGCTCC
10681    AAGTAAACAT TCCAAGTAAG CGTCTTAACA CCTAACCCCA AAGGCTCTTT TAAGAGCCAC
10741    CCAGATACCC ACTAAAGAG CTGTGGCCAG ACGCCAAATT TTATTTGGCG GCGGAGGGGT
10801    ATTAGAATGT AGGAACTGGA GAGGGGTGGG GACAAGTGTT GCAGCTTAGA GAGGGACAAA
10861    GGGTCCTGAA CCCGAAAGAA GCCAGCCATT AAAAATGGGG TTGGGGTCAA TTCGTTGTGC
10921    TTAAATTTAA AATGGGGACA AGCGGCCATT TTGCTAACTC GGCGTTCCCG GAAGAAACCG
10981    CAGGCTCGCT TAGGTTTCAG ACCCAGCTGT CTGTCCCTGT CTACGTCGCC AGGATCAACG
11041    GTTGCCGTAA TGTCATAATT TCGCCACCAG CTTCTAGCCA ATAGGCTGTC CTGTCATTTT
11101    AAATATTAAC CAATCGAGGG AAAGCTGTTT TGAGACTCTG ATTTACATAG CGGACCGGAG
11161    TGGGAACCTG GGCAGTAACT GCCTAAGGAA GGACTCCCCC TCTGTTTTCG TGGCGCACAC
11221    CTTCGTAGTA TACTGAAGGG TGTGTCTCCT GGGTTTCCAA CTGCCCCGGT AATAGTCTTT
11281    TAACCTAATA TGCGTCAGTT TTGATAACAA CACTAAGGCA GTACAGAACT AAAGATGTAA
11341    GCACTGCGCC AGATGTTGCT TCATACATCT TATTCTATTC AACTGGTTTA TTCAAGATTC
11401    AAATCAAATC AAATTTTGCT TGAATCCCAG TGCTCAGTCA GCCATAAATG GTGTGTTGCC
11461    TGATTGAAAC TTAAAATCTC CGTAGGGGGC TTGTAACATG CAGAAAAGTT TGAAAGTTGC
11521    TTTAGGAGAA GCCAACTCTT AACTGCTGGG TAAATTGACA AGCCTTCGAA CACTGAACTG
11581    AAGGCCAGTA AGGACTAGGC GCTGGGTGGG GGAGAATGAA GAGGAGACGT CATTAAACTT
11641    AGCACATACA CTGTGTCTCC TAGAGGACTC TCCCTTCCTA GACAACTGCA GGCCGCTTTG
11701    TGGCCTGGGA AATTCCACAT TCCCTTAAGT ATTTTACTCA TGGTCTTTTC CAGGTAAAGA
11761    TTTTAAGATG AAGGGTTAGA CGTAGTCTAC CTATCTTTTT ATTCAAGTCT AGAACACGTT
11821    TTTAGCACCT AGAAGTTTGC TTTCTCCATT AAAAACCGGG AATATACAAT AAATAAAATT
11881    AGTGTTAAAG CAGATTTTTA CAAACTTAAA TACCATGTAA TTTAGGTTAC AGTTACTTAA
11941    CATAAGGACT GTGTGATCTT AAATCTGCAA TTTCTTTCAC ACCTGGGAAA TAAACTAAGG
12001    CCTGTCTTTG GTGCCAGACA AGGCCTTATA CTTGAACACT GCTGTGCAAT CACAGGCTGC
12061    CTTGCCTAGA TAACTTATCT GAGAAATTCT GATGAGAAAT GAAATTTCCA GAGTCCCTCA
12121    CAAGTAAATT TTTTTTTCTT TTTTTTTTTT TTTGAGACGA GTTTCTCTC TTGTTTCCCA
12181    GGCTGGAGTG CAATGGCGCG ATCTTGGCTC ACAGCAACCT CCGCCTCCCG GGTTCAAGCC
12241    ATTCTCCTGC CTCAGCCTCC GGAGTAGCTG GGATTACAGG CATGCGCCAC GACACCCTGG
12301    CTAATTTTGT ATTTTTAGTA GAGACGAGGT TTCTCCATGT CGGTCAGGCT GGTCTCGAAC
12361    TCCGGACATC AGGTGATCTG CCCGCCTTGG CCTCCCAAAG TCCTGGATTA CAGGCTTGAG
12421    CCACCGCGCC GGGCCTAAAT GGTTTTTTTT TTTTCTATGC CTCTAATGGA CCTGGTCACT
12481    TATTCCCATT CAGACTGACC GCTCTCCTAC CTGCCAACTA ACTAATCAGT GTAACCAAAA
12541    TCTGCAAACA AAATTCAGTA TTCTTTCCCC GCCTTTTCCC CTTTCTCTTA CATAGATTAT
12601    GTTTTTGCCT GTGTTAGATG AAATAATTCT ATTGCTTGTT CTCTCTTCTG TACAAGTACC
12661    CAGTAAGCAA ATTATTAACT TCTTGGTCAT TTATTTCTGA ATTTTCCACC AAGACAGTGT
12721    TTATGTGAGT CATACAATAA GAACCAACAG AAATGTGTGT CTTGGAAACA GGTTGTCTAT
12781    CCCTGGACCC TTTGAGTTTT CTGTTCACTT TCCTTTGGCT TTTGCATGCT AAAAGTTTAT
12841    CGTCCGCGTT TGTTTGTTTT GGTTATTCTA ATTGGACTTG GCTGATTGGT TGCATATTGG
12901    TGGCAGTAGT AGAATTTGAA TTCTGGTTTT CTGGTCACAT CATTAAGTGA TTAGTCAGTG
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12961   GAGAGGACAG GAAATCTGGT TTATTTATTA ACCTTTTTTT GGGGTGTTTT TGTTTGAAGA
13021   TGTTGATATT CTCTGTGAGG ACACAGGGTT AGAGTTGGTG TTTTTCTTTC TGACTTTACA
13081   TGGGATTTGA TGTTTTGTGC TTGTATGCCT CTTTCCACCT TCCAAAACTT GTCTTTTTTG
13141   AGTCCAAATA GTTGTCGATA TCTGCAAAAC CAGTATTCCT GTGTTAAGAT GATATGAATA
13201   TAAAATGGCT GCCCTGTTAT AACTTTTGAC TTTAAGAAAG TGTTAGGACT AACAGGAGAC
13261   AAAAAGGAAA TCAAGGAAAC CAAATGTCTG GTCTCAATAA CTGCTATGGC AGAGGCTCTA
13321   CAGCTTATTA TTAATTTTAG TAATTTCACA TTATTGCCCC TTCACGTTCT TTAAGTAAGG
13381   TTAGAGGACA GAAGAAACAT AATGTTGTTA CAAATTGGAC TATTGAGTCA GGAAAAAAAA
13441   AGAGTGCTTT CAATATCTGA ATAAAACAAA GATTTAATAT TTTCTAAACC TTAACGAGTT
13501   TATTGTAAGG GATGTGATGC TGGAAACTAG GAAACTAGAA TTTTCTTCTA AACTGAGAAT
13561   CAGAATTATT CATATTCTCA GCAGTGGTGC CACCTGAGGG ACTTCTGATC TTAATTACAT
13621   ACTTTTATTT CTTTAACTGA TCAACATGCT AAATAGATAA CCTATGGCTC TGTTTTTACC
13681   CACTTTAAAT TCTGTTCTAT TAGCACGGTT AGCTTTCCTA ATTGGCAATA AGATTGAGAC
13741   TATCTTTTTT TTTTTTTTGA GACAGAATTT TGCTCTGTGG CCCAGGCTGG GGTGCAGTGG
13801   CACAATCTCG GCTCACTGCA ACCTCTGCCT CCAGGGTTCT AGCAATTTTC CTGCCTCAGC
13861   CTCCCCAGTA GCTGGGATTA CAGGTGCACC ACCACGCCTG GCTAATTTGT GCATTTTTAG
13921   TAGAGATGGG GTTTCGCCAT GTTGGCCAAA CTGGTCTCGA ACTCAGGTGA TCCACCTCGG
13981   CCTCCCAAAG TGATGAGATT ACAGGCGTGA GCCACCGTGC CCAGAAAAGA CTATCTTATT
14041   TTATGAATTT AAATAATTGT GAAATTATCC ACTTAAGGGA ATTAATAAAT TATAATGTAA
14101   TCTTAAATTT TAGTTGGCTT ACATAAAGAC TTAAAATACA TCAATTTAAA TAAAAACTCA
14161   TTTGTCTAAA AAAAAATCAA AAATTTTCCT TGTGCTTTAA ATGTGCTACC TCTTTAAGTT
14221   CTAATTAAGA GAAAAAAAGT TTAACTGTGA GTTTCATTAG TGGTCTTAGT TAACAGCTTA
14281   AAGTATTTTG TAAAAAAAAT ACTTCACAAT TTTTAAATAA CTTAAAAATA TTAATACCTC
14341   TTTTATTAGG TTTTTTTAAT AAGGAAAATA TATAATACAT CTAATCAAGA TTATTTTTTG
14401   GACAAATTGG CTTAATAATT TCATTTTAAA AATGGCTTCT TTATTCTTAT ACTGTAAAAA
14461   TAATATTAGC AGAATATTAT AGTATACACA AGTTTAGGGT TCATATTCTA AAAAACAAAA
14521   ACAAAAGCTA ATTTAACTTG CATTTACTAA ATTTCTTCCA CTAGTTGTAC TGGTTACATG
14581   AGTTAACATC ACTTTATTTA TTATTCTAAA ATTGTAAATT ATTCATTGAA CCAAATTAAA
14641   TGATAATAGA TAATGTCATT TTTAAAAATG GAATTAAATT TTATGTTACT AATTATAAGG
14701   ATTCAATGTG TGAGCTTAAG TACTGAGTTC ACAGTGTATG ATAACTTTAA GAATTTAGGT
14761   GAATATTATT AAATTGAGTA AATTAATTCT CAATCTTTGG ATACCTGGAC AATTTCTAAA
14821   TTGGAGGGTA CAAAATACAA ATCACAAGAA ACAGTGTAGT TTTATGCAAA TAACATTTTT
14881   ACACAGTTTA GAATAACCAT TGATAAACAG ATAAGAGAAC ATATGATTGC CTTAGAATAG
14941   ATACTGTTGC TTTCGCCACT TTAGATTTGT AAATCATGTA CTGTATACGT GTGGGCGTAG
15001   AGGACCATGC AGGTTTTGGA TGACTGCCTC TGTTTTCGTC ATGCCTATGC GGGAACACAA
15061   TTGCCTGCTT TGTTTAAGGG CTATGGTTAA TCCAAACAGC TCTGACTCTA TCAAGTACTA
15121   TAGCTACAGA GAAACACAAG TAAGCATTCG AGATAATGAC TACCTTGAGC CTTTACTTAT
15181   TTAAAAAGTT GTTACTGTTT GTTAATGTGG TACATTCAAT TTACTATGGA TTGTCACTCT
15241   AAAATAAGAC TTCAATCTTT TTCTTATTTT TATATAGCCA TGATTTATAT TCATATCTTA
15301   ATGTAATAAC CAATCTTCTC TGACAACATT ATAACAATGC TGGAACCTCC ATTTTCAGTA
15361   CTTCAAACAA CAAATACTGC TTTTATACTT CAGAGCAGAT GGATATGTGC TTCCCAGTGT
15421   AAACACATTT GGAATCTCAC TGAGAAATAC ACTATCACTA AAAATACAGT TCTGAGATTC
15481   ATTAAAAGAC CTCCAGAATT CTGGAAGTAG GAAGTTTCCT CTTCAAAGTC TACAGAGGAA
15541   GACGAGGTCT GAAATAGACA GCTTCTTCCT TCTTTTACCT GTGGTATTAT TCTGTTTTGT
15601   CCTTTTCTCC ATTATCTGTC TTTCCAGTGA TGAAATTTTG ATCTGGCCCT CCCAAGTATT
15661   AAAAAACAAG CAAATAAACA AATCTCAGTT ATATTTACT AAGATATTGG CATGCTAACT
15721   TTTTGCAGGT TTGTAACAAG GACCTTTATA ACTTGACTAA AAGTTCCTAA ATAAGAATAT
15781   TTACTAGAAA ATTTATTTCT GCCTGTGGCC CACATTTGAG TCAAAATAAT CAATTAGGAA
15841   AAATGAACTT GTTTAACTAA AGTTGGCCAA ACTGATCTTT GAGACCTATT CATCTAAGAC
15901   AAGCCAATTA AATTCTTGGA GACAATTTGT ACTTTAAGGA ATTCTTATAA TATTTGTAAT
15961   TACCCTCATA ACTTTTTTTT TGCCCTACTT CTGTGCTTCT CTAATATGCA GATTATTAAA
16021   TGTTGTTACA AAGCCATTGT CAAAAAAACA AAAACAAAA AACTAAACAA ACTCACATGG
16081   TTAGACTTGC TCCTTTATGA GATATTTTTA CCAAAAATGG AGGAGTTGAA AAACTCTGGT
16141   GCCAGAAATC GTGAAGACAT GGCCTACCTA ACTTGGAAAT GTTGGTTGTC AGTGGAAAAT
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16201   ACTACACAGA GATAGCCATA GTGCTGCACA GCCAATCTTA AGTGTTTCTA GAGAATCACT
16261   AATTGTTTCT AGAGAATCAC TAATTGTTTT CTTTTAACAT TCTTGGTTTA TACAAGAAGA
16321   GAGTATCCAT ACTAAACTCT TTTCTACTGA AAATAATGTG CAAACATAAC ATCCTATTCC
16381   TAGACAGTTT GTAGTTTTTT TCTCCCATTT CTATTTTATA AATCATCTTT TTAAAATACT
16441   TTGTTGAGTG AAATCAGTCC ATTGCTTGAT ATACCTTGAG CACAAGTAAA TAGTATGCCA
16501   AAAATTAAAT GTCTTTCAGT CACAGTTTGA CAAACTCAAC TACCCTGAGC CTATAGAGTG
16561   GTAATAATTG CCCTACTCAT AAAGATGGGG TGAAGATTAA ATGAAATAGC ACCTATAGAA
16621   CACTAGTTCC AGACGTGGTA TCATGCTAGT AAAATGGCTG CACAGCACTG CTCAATGATG
16681   ACAAAAGTG AAGCTTCTGG AGACAGACTC CAAGTTTGAC TCCCAGATCA CCACATATAA
16741   GATGTGGGAC TCTGAGGCAG GTCATTTAAT CTCTCTGTGC ATTAGTATCC TTCTCTATAC
16801   CTTTACAGTG ATGGTAATAG CACCTACCTT CTAGAAGTAT GTGAAGATTA AAGATCCTTA
16861   ATGCATATAA ACCACTGTGT TTACTGCTGT TTGACAAATT TTATTTATAA CCATCTTTAC
16921   GCTCCTAAAA GGACTTGAAG CAGCTTATGA CTGAAGACTT TGGTAGGAGT TGGCCTTCTA
16981   TAAATTATAA GAATTTCATA AATTATTTGA TATGAAAATG CCAGTTGATC ATAGTATGTT
17041   TACCGGGGTC CAACAGGTTG AGAAAAAATA CACTTTTTTT CCCTGAACAT ATGAAATTAG
17101   CTCTCTAGGC ATATTCCTAA GGACTTAAAG AATGATAACT ATCATTTCTC TTAAATCTTC
17161   CAGATTTGGA AGGATATATA TATTCAGCAC ATTGACAGAC AATCCCAGTA GTCCTAAATT
17221   AAAAGACATT AAAAATTAGT GAAACTTTTC CTACCTTTAG CCTGTGTAAT CCTGGATGAC
17281   CAAGCATAAA ATTAAATTGA GTAGAGTATA CCACTGTAAC ATTTCCTGAA AGGTATTCTA
17341   GGCTCTGAGT AATTTCTTTG GGGTCTGAAG ATCAGTTTGA CATATCCTCA AGTATCATGA
17401   GTTCATTATA ATTAAGAAAA AGGGAGTAAA TCTGGAGAAT GAGCCACTTT CTTACTACTC
17461   CTTGACCTCA GTTCTTTTTT TCAGAGACAG GGTCTCACTT TGTTGCCCAG CTGCCAGGC
17521   TGGAGTGTAG TGGCGCAATC GCATCTCATT GTAACCTCCA CCTTCTGGGC TGAAGCCATC
17581   CTCCTGCCTC AGCATCCTGA GTATCTGGAA CCACAGCAGG TGCACACCAC CATGCCAAGC
17641   TAATTTTTTA AAAAGTTTTT TGTAGAGATG GGGTCTTACT ATGTTGCCCA GGCTGGTCTC
17701   AAACTCCTGG GCTTAAGTGA TCCTCCTGCC TCAGCCTCCC AAATTGTTGG GATTACTAGT
17761   GTGAGTCACT GTACCCCGCC CCACTTCAGT TCTGAGGAGG AAAAAATATG TAATAATAAT
17821   GGGACTTTGG TTTGCTGATT TAAAGATTCA TGTAACCTTA TCATCCAATG CGCAATTTGT
17881   AGAATAATTA ATAGAGACAT CTGGTCTCAT GTTTCTACAG TTGCTCATGC CTTGATAGTA
17941   GATCTCCTTG CTGCTGGCTC AGAAGGGTAA AAGAGCAGAA ATGATGGGGC TTCTCTCATT
18001   CTATGAGGAA ATAGACCTAT GTAGAGGAGG CTACCTGTGG TAAAACCTTA TCCTCATCAC
18061   TTAAAATTCT AGGCTTATTC TCTGACCATA TCAAGTTTTC AAATGGTAAA AGAATTGGAT
18121   TCAAGAGAAA TATGAATAAA CTTTTGTTTT CACTTTTCTC CCTCCTCTCC CCCCATTCTC
18181   CCTTCCTTTA TTTTCTTGTC CTTAGTTTTC TTTTCACTTT TTGTCTACT ATTATTTGCC
18241   CAAACTCAAC TGTAGGCTAG AACAAAAAA AATTGAAAAT TAAAATGTGC CCCTTTTGTT
18301   GTTAGACTTG CTTAAACAAT TGGGGTAATG AACCTTGGAC ACTAGATTTT AAAACACACA
18361   CATTTGAGCT TCAGTGCACT GAAATAAATA TATTTTTAAC AATTAAAAAA TAAAATTGCA
18421   TGTTTAAAAA ATCTGCAGAG AACAATACAC GTTGTGAGAT CTTGAATGGA AGGAAAACTG
18481   CTAGCCTCAA GAGTGGATCA AAGATGCTCA GCAGGCAACA GAGTAAGAGC ATGTTGGAGG
18541   GTTTAGAGAG TGTGCTCAGG GTTCTAGGCT CTAAAAATCA GACAGTCCCC ACGGCCTGGC
18601   CTTCGTCGCT GTATCTTCTT TATGAAAAAC ACTAAGTCTT TTTCCTCACT GGATAAATTT
18661   TTATCCTTCA AGTTTAGATC AAATGGAACT TTAGGACACT GACTAGGTTA CATTCATCTT
18721   TTAAGAGCGT ACAGACATTC AAGGGCTAGA GGATGTGGGT TTACTGCACA GGCTCATTAT
18781   CCAACAGCTG TGCTACCTGG GAAACTTAAC CTCTCTGTGC CTTAATTTCC TCATCTATAA
18841   CGCAGGGAGA ATGACAGTAG GTATCTCATA AGGTTGTTGG AACAACTAAA TGCATTGGTA
18901   TCTATTGTGT AAAGTGCTTA AAACACTGCC TGGCACAGAG CAAACATCCA GTGAACTTTA
18961   GCCATCATCA TTATCATTGT TCTCAGAGTC AAATACAATA TCTCATATCT GATAAATTAC
19021   AGAAGTGAAT CAATCACTCT CTCTCTTTTC TCCAGGGGGA GACAACAGCT TTTAGACATA
19081   TCTTTTCCAA CAGTCGTCAC TGCTGGACAC TGTTTCATCT TGCAAATAAA CCAATGAAAA
19141   TGAGTGATCC TAGAAGAAGA TAAATGGAGG TATTTTGAAC AATCAAAGAA GGACAAATGA
19201   ACACCTGGCT GAGAAAAATT AGCTCTTTTT TCTATGCATA AAACTATTAA AATATTCTTC
19261   ATAGAAATTT ATGACACAGG AAACATAAAG ACAAAATTAA AATAACTCCT AGTATCTCCT
19321   ATTCTTTTTA TATGTATATT ATATATACTC ATATTCATAT ATACATATAT CTCACATCAT
19381   GTATCATATA TAAAATAAAT TTAGGTGTCA TGATATATAT TTAGATAAAT ATACTTAGAA
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19441   ACTTTTTTAT GGATGTATAA TTTATGGATA TATTGATAAT TATGTATTTG TTATTGACTA
19501   CTTCAATTGA TTCCCATTTT TATGCATTAT ATTATAGATT ATATAGCTCA CACATCTTTG
19561   TACATAAATC TTTGTTCAAA TATTATTTCC TAAGGATAGA CTTCATGAAG TGGAAATACT
19621   AAATCAAAAG TGAAAAACAT TTTCTAAGGT TCTTAACATA TACATTGCCA AATTGCTATT
19681   CAGGATCATA CCAATTTATA ATCCCAAAAT AAATATGAAAA TTCCTGTTTT ATAGCACTCA
19741   TATTTACAAT AAATTTTAAA AATCACTGTT AACCTAATAG TCCTTCAAAA GAAAAAAAAA
19801   TTGAAATTAC ATTATTTTAA TGACTCTATT AGTGAGGGTC ATTCTTCCCA TGTTTCTTGT
19861   TAGCCATGAC CCTATAAGAA ATAAACTGCA CTGCAAAATG ATAAACATGA TATCAATCAT
19921   TACATGGGAA GGCACTATAT AAAGAATAAT ACCTTAGGTT AAGGCCACAT AAATATTTAT
19981   CAGGTGCCTT TTCTGCGGAG GACTCTGAAG GGATACTAAA CTGCATTTAG CTGCATGCAA
20041   CTGAAATTAC TTTTACCTAC ATTGTCTCTT ATAAACATTA TAACTACTCT TTGAGAAAGT
20101   GTTTACTATG GACTGAATTG TCTCCCCATC CCCCCAAATT CATATATTGA AGCCATAAAC
20161   CCCAATATGA CTCTATTCCT AGACAGGACT TATAAGAGGT AATTAAGGTT AAATGAGGTC
20221   ATTAGGATGG GTTCCTAACT GGATAGGATT GGTGGCCTTA TAAGAAGAGG AAGATTCTGC
20281   ACTTGGTCTT CCAAATTAAA TAATTTATTT AAAAGAAAAA AAAAAAAAGA GGAAGAGAGG
20341   GAGCTCTGCA CATATACTGA GGAAAGGCTA TGTGAGCTCT CACAGTGAGA AGGTAGCACT
20401   CTACAAGCCA GCAAGAGAGC CCTCACCAGA ATCCAGCCAT GCTATACCCT GCTCTGAGAC
20461   TTCCAGCCTC CAGAACTGTG ATAAAATTTT GTTGTTTAAA CCACACAATC TATGGTATTT
20521   TTTTATGGCA GCCCAAGCCA ACAAAGACAG CATCATTGCT GTCACTTACA GACAAGAAAA
20581   CTAAGACTAG GAGAGAGAAA AGTTAAACTT GTCCAAGGTC ACAAAAGCCA GAAACAAGTG
20641   AGGTGAGAAG TTGACCTTGT TCTCCTCAAT CCAAGGCCAG GACTCCTCCA CTCCACATGT
20701   AGATAGCCAC CTCACAGTCA ACAGCCAAAT GTCCACACCC CAGAGTCAGC ATTAGACCAA
20761   GATGTCTTAC CAGGAGACAA ATGCCTCATC TTGAATAAAT ATGTTCTAAC AACTTACCCA
20821   TGTAAAACAT TGAATCTCAT GAGAAACAAA AATGCAAAGT ATGTAGAAAA CTATGTTTAC
20881   CACTTAACTG ACAGTGATAA AAAGCTTAAT GATATCCTTA TAGTCTTGGA GGGGTTTGTA
20941   TATGTGGTGA AACAGGTGCT CACGCACTGC TGATAGACTG TAAATTGGTC CTAGAGAGAA
21001   AAATAAATAA ACTGGAAGGA GTTATGCTGT ATGTTTACTT TTTTATGGA AACATATGAT
21061   ATACCTGGAA ATTCGATTGG CCATGCATCT ATTTCTTCAA TGGGTATGCA CAGTTGAGCT
21121   GTTCCCATGC ACCAGGCACT GTAATGGGAC AACTGCACAT GACAGTCAAA AATCTCAGTC
21181   TCATGAAGTC GACATGCTCA TGGAGAGGTG CTACCCACTA AACTAATATT TGTATATCAA
21241   TTATGGATAC ATTGGGCCAC ATTTACAGAA ATTCACTTAC AGTGGGTTAC CAGAAGGGAT
21301   TTTTTTTCTT GATTGGCAAG AAGGCTAGGC TGTTTTGTTG GGGGCTGGCA GGAGCTGTCT
21361   AGGCTGCCCA AGTATGCAGG TCTCTTCTAT CATCCTGTGT TAACCATCTT CCATGTATCT
21421   TTCAACCTCA TGGTCATCTG CAGCATGTCT AGGGGTCATA TCTATGTTCC ATGCAGGAAA
21481   AAAGGGTAAA GGGAAAGGGA AGTAGGCATG TACCATTTTA ATGCACACCT TGGTTTTCAG
21541   AAAATTTAAG AAGAAAGACT TTCTGCTTTT CTCTGACTAT TCTGTATTCT GGATTACAAC
21601   GCAACAGAAA CGTCACCTTA AATTCTAATG TTTTTCTCTC CTTGCTTTCA AAAACTGACT
21661   CATTAACCTC CACGTGGCTT GGAAAAATTA TTTCAGTCAT CCAGTAATGA GCTGTTCATA
21721   GAAATGTTTT GGACATCAAG TCTGTGTTGT TAGCATTATA CATGTTAAGC ATTGAATAAA
21781   AAACAACATG ATGTGGGTAC ATTTCTTTAC TTACATATAA GTACTTATAT ACTTATAGCT
21841   GAAAGAGAG GTTGAAATGT CAGGTGGAAC AGAAATAAGA TTACCTAGAT GTTTCTCCTA
21901   TGGGTGATTT TCAGCTATGC TGATCTTTCT TCTGGGTCAG GTACTCCCAG AACTTCCTAA
21961   TTAAATGGTG GCCCTGATCT TAGTTCCTCT CTCCTCTTAG ACATTTTCCA GGACTACAGA
22021   AGATGTGCAG TTTATAAATG AGTAGCAGAA ACCTACTGAA CAAATTATTC AGGCTCATCT
22081   GAACAGAGAG GACACCTTCT CTGCTATACT CTCTCAGTGA TTTCCCTGCC TTGGGGTCAA
22141   TTATTGTCTT GGACATTGAT TTAAGCACAT AATAATTGTT GTCATTGCTT ATGTTTGGAT
22201   TTCATCTCCC AAAATAGATG GTAAATTCTT TAGTTTAGAG ACCAAGTAAT ACTTACAAAA
22261   AAATTTGTG TGTGTGTGT TGTTTTTTCT GTGTCTCTCA GCCCTGTAAT AGCATCGTAC
22321   TTACACTTGT TAGATTTTTA GAGACAACTT TTACAAAACA TGGAATTATC TACATACCCT
22381   TTCTACAAAA CAGACAAATT AAATACTCAG TAGTTGAACC AAAAAAAGCA GTTCAAATAA
22441   AATACTTGAA AATGAAGAAA TCATTTGAAC AGAGTTAAAG TTAATCGTAA AATAATGTCT
22501   GTAAAAATTA TTGCCAATCA AAATATAAAGT TCAAAAATAG TGCTTGAAAA AGGAAGAATC
22561   ATATGAAAAG GGACTACTCA TTTTAAAAAT GTTAGATATC AGGAAAAGCC AAGAAGTGAG
22621   TATGGTAAGA GTGCTGTCAA GTGAAACCCT GCTAATCTCA CTGAACATGT AAAAATCTGT
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22681   AGATGCCTTT ATTTTATTCA CTCACACACA TATGTAGAAA GAGAAATATA TGGTAAACAT
22741   TAAAAAAAAC AAATTAGAAT GTAAAATTAA TACTTTAAAA AATGGGCTGT ATACTTTTCT
22801   TATCACCGGA GATAAGAATT TATTATTTTT AAAATAAAGT TATTTTCTCT GTGACTGTTT
22861   CCATGACTTT GCTACTTAGA AGTTAGAGAT GCCAAAGTTT ATCTAAGAAA ATGTTTATGG
22921   AAATATTATT TCAATAATGA ATGTTTAGAA GACTGAATTT CCTGACTGGG CACAGTGGCT
22981   CATGCCTGTA ATCCCAGCAC TTTGAGAGGC TGAAGAAGGA GGATCGCTTG AGTCCGGGAG
23041   TTCAAGAGCA TCCTGGGCAA CACAGCGAGA CCCTGCAGCA AGTAAAAAG AAAAAGAAT
23101   TGAAAAGGA AGACTGAATT TCCTTTGGGC AAGTCATGTG ACATTCCTGT GCCTCAGTTT
23161   CTTCATCTAT AAAGTTAATT CCTACATTTT TGGGGAAGGG AGAGAAAAAC TTAGGATAGT
23221   GACTGGCACA GAAGAAGCAC TATATACTAT ATATATGTGG ATATCATTTG TTTTTATGGT
23281   ACCATTTTAG CTATCTAATG CAAAATATGA ATCTTTTTT TCTGGGTCTT AAATTATGGA
23341   ATGTAAGAAT TTTCTAAATT CTCTAATTCT GTGTTAGTTT TAAAGCAATG GAGTAACGTA
23401   TCTGTCAACT TGTAAATATA AGGATCAACC TGATCCACAA TTTGACCCCT AGCCACTAAT
23461   ATTTAATAGT ACAACACTCA GAAATTATCA AAGGTCAGAG AAGCCAAACA AATGTAAAAA
23521   CATACAGGTG CTCAGAAAGA TGCACCTGTA ATCTCTCTAA GGAGAAATAT TTTCCAAACT
23581   GAGTGACACG GTGCTTTAGT GAGTTGTGGA ATCAATCTCA TGATTTCCAA CCTAGTGTTC
23641   TTTTAAAAAT GAACTAGTCC ACAGTAGAAT ATACTAAAGT GCTGGTGCTT AAGATAGTAT
23701   TGTTTTCTGG AAAAAAAAAA AAAATTTTTT TTTTTTGAGA CAGGGTCTCG CTCTTGCCCA
23761   GGCTGAAGTG CAGTGGCACA ATCATGCTCA CTGCAGCCTT GACCTCCTGG GCCCAAGTGA
23821   TTCTCCCACC TCAGCCTTTT GAGTAACTGG GACCACAGGT ACGTGCCACC ACACCCGGGT
23881   AATTTTTTAA TTGTAGAGAC AGGGTCTTGC TATGTGCTTA GGCTGGCCTT GTGAACTCCT
23941   GGGCTCTAGT GATCCACTAG CCTCAGCCTC CCAAATTTAT GGGATTATAG GCATGAGCCA
24001   CCCTACCTGG CCTGTTCCCT GAATTTTTTT TTCTTTCAGG TGTTTGTGCA TATGTGTGTG
24061   TGTATGGGTA TAACAGAGAG ACAGAGAGAA AGAAACTTTT CTATCACACT TTGCAATCAG
24121   AAGTTTGAAG TCTTATCTTT TGGCTTTTGT TTCAGAAATA TTTCAAATGT AGACTCTCTC
24181   CTTTACCACA CTGTCCCCTT AGGCAAGGTC TTTGCCATTC TTCTGAGACT ATTGCAACAG
24241   ACTCCCAACT TCTGACTGTG GGCCCTTCTC AAAAATGATT GTTTATGCAA TAAATCTAAA
24301   CCCAAGACAA CTACAACAAT ACAACAAATT CTCTGCTTAA AAACTTCCAA TGTCTGCCGG
24361   GCGCGGCGGC TCACGCATGT ATTCCCAGCA CTTTGGGAGGC AGAGGCGGGC AGATCACTTG
24421   AGGTGGGGAG TTCGAGACTA GCCTGGCCAA CATGATGAAA CCCCATCTCT ACTAAAAATA
24481   CAAAAAATTA GCCAGGCATG GTGGTGGGCG CCTATAATCC CAGCTAATTG GGAGGCTGAG
24541   GCAGGAGAAT TGCCTGAACC TGGGAGGTGG AGGTTGCACT GAGCCAAGAT CACACCATTG
24601   CACTCCAGCC TGGGCAACAA GAGCAAAACT CTGTCTCAAA CCAAACCAAA ACAAAACTTC
24661   TAATATCTAC CAAATGTTTC ACACAAGTAT TTGGGGATCT TCACAAATGG CCCTTATGGA
24721   GTTTTCCTTT GCTGAGACCC TATGCTCTGG CCACACTAAA CTCATTCAGC ATCCCAGAAA
24781   GGCCTCAGCC TTTGTGAGCA AGCTCTTATC TCCAGGCCTC TCACAAAGAC CTGTTCCAGT
24841   AGAAGCTCAG GGGAGCACAC TGGACATTAT TCCAACAACC CTTTCCCCAC AGCTATGCAG
24901   CCAAATCTGC CAGCTCAGTT AATTAATTAA GCAATTCAGA GATGAGGGTC TGCCCAGGCT
24961   GGAGTGCAGT AGCTGCGACC TCAAGCTCCT GGGCTCTAAG TGATCCTCTT CAGTCTACCC
25021   AGAAGCTGGG ACTGCAGGCA TGTGCCACCA CACCCAGCTA ATTTTTTTTT TTTTCAGTAG
25081   GGACCAGGCC AACCTAGTCT TGAACTCCTG GCCTCCAGCC TTCCGAAGTG CTGTAATTAC
25141   AGGCATGAAT CACTGCGCCC AGCCAACCCG CCCAGTCTTG TTAGACATGG GGTCTGTAGT
25201   TTCTAGTAGG TTCTTGAGTC TAGGGTTCCT ACCTCATGTT TTATAGTTAA TTTAGGGGAG
25261   GGACTGTGTC TGTTTATCTG GGGATGTAGG GGTGGGCAGG GGGATAGAGG GGACTTCAAT
25321   TAATGAAACC AGAAGCAAAA CTCAGTTGAG GACACCGGTC ATGAGAGTGG CCTGATTATG
25381   GCCAATCTTA CATAATGTGT GAGATCTTGA TATTACCCCA TCCTTGAGAG TCCTCTATAA
25441   AGCTACAGGG ACTTGGGAGC ACCTTTAATT ACAGACAACC CATGTTCCTG TGGATTATGA
25501   TTTATTAGAT TGCACATGCC TAAATAAAGA CATCCTCTGC AGTCTTTTGA CAATTCTATA
25561   AGCATCTTCT GACTCCGCAA TTAGACAGCT AAGAGATCTG TGTTACTTCC CTCACATATA
25621   TAAATAATTT TAAATAAAAA TCATGGCGTG AATAATTTCT TTCCTCTACC GATTTGAAGC
25681   TATCCATTTG GAAGACCACT CTGAAGAGAT GAAATAAGTC TTCTGCCAAA GATTACTTAT
25741   TAATTTACAA GGAAAAGGGG AAGTTTTGTT CCTCTCCGTG AATTTGATTG AAAATCGAGG
25801   GCTTTCTCGA ATAGTTTTGG CATCCAGGGT CATTTTTCAT TAAAAAGAGA AAAGTCATGT
25861   CAAATATGAA TTTCCGCAGA TTATTCAGCA CTAGACCCTG GGAGATTCTG TAAAGAGGGG
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25921   TTTTGTTATA CTCAACTTTT CCGGGTAAAA CAAACACAAA TACTCCTCCT CCAAGGGGCG
25981   GGGGCGGTGC CTAGGTGATG CACCAATCAC AGCGCGCCCT ACCCTATATA AGGCCCCGAG
26041   GCCGCCCGGG TGTTTCATGC TTTTCGCTGG TTATTACATC TTGCGTTTCT CTGTTGTTAT
26101   GTCTGAAACC GTGCCTGCAG CTTCTGCCAG TGCTGGTCTA GCCGCTATGG AGAAACTTCC
26161   AACCAAGAAG CGAGGGAGGA AGCCGGCTGG CTTGATAAGT GCAAGTCGCA AAGTGCCGAA
26221   CCTCTCTGTG TCCAAGTTGA TCACCGAGGC CCTTTCAGTG TCACAGGAAC GAGTAGGTAT
26281   GTCTTTGGTT GCGCTCAAGA AGGCATTGGC CGCTGCTGGC TACGACGTAG AGAAGAATAA
26341   CAGCCGCATC AAACTGTCCC TCAAGAGCTT AGTGAACAAG GGAATCCTGG TGCAAACCAG
26401   GGGTACTGGT GCTTCCGGTT CCTTTAAGCT TAGTAAGAAG GTGATTCCTA AATCTACCAG
26461   AAGCAAGGCT AAAAAGTCAG TTTCTGCCAA GACCAAGAAG CTGGTTTTAT CCAGGGACTC
26521   CAAGTCACCA AAGACTGCTA AAACCAATAA GAGAGCCAAG AAGCCGAGAG CGACAACTCC
26581   TAAAACTGTT AGGAGCGGGA GAAAGGCTAA AGGAGCCAAG GGTAAGCAAA AGCAGAAGAG
26641   CCCAGTGAAG GCAAGGGCTT CGAAGTCAAA ATTGACCCAA CATCATGAAG TTAATGTTAG
26701   AAAGGCCACA TCTAAGAAGT AAAGAGCTTT CCGGGAGGCC AATTTGGAAA GAACCCAAAG
26761   GCTCTTTTAA GAGCCACCCA CATTATTTTA AGATGGCGTA ACACTGGAAA CAAGTTTCTG
26821   TGACAGTTAT CTATAGGTTT AAGTTGTGAT GCAGCTGAGT TGAAAAGGCT TGAGATTGGA
26881   GAATTAATTC AGGCCAGGCT TCAAGACCAT CCTGGGCAAC ATAGCCAGAC TACCATCTAT
26941   ACCAGGGGTC CTCATTCCCC CGGCCACCGA CCGGTAACCG GTCCCTGTCC ATGGCACGTT
27001   ATGAATTGAG CCGCACAGCT GAGGGGTGAG CGAACATTAA CCAACTGAGC TCCACCGCCT
27061   GTCAGGTTAG CTGCAGCATT AGATAGATTC TCATAAGCTC AAACTGTATT GTGAATGGCA
27121   CATGCAAGGG ATCTAGGTTT CAGGCTCCTT GTGACAATCT AATGCCTGAT GATCTGAGGT
27181   TGGAGCAGTT TTAGTCCGGA AATCATTGCT CCCAGCCCCT GCACCCCTG TCCGTGGTA
27241   TAATTGTCTT ACACAAAACG GTCTCTTGTG TCAAAAAGGT TGGAGACTAC TGGTTTTACA
27301   AAAAAGTAAA TTAGTCAAGC ATGGTTGGCA CGCTCCCTTA GTCCCTGCAC CCAGGCGTTT
27361   AAGGATACAG TGAGCTATGA TGGTGCTACC TCACTCCAGC CTGGGTGACA GCGAGTCAGA
27421   CGTTGTCTCA AAACTTAAAA AAAAAAAAG TTAAAACAGA AAAAGGGCTT CTTGTCAGAG
27481   ACTGCCGTAT ATCTAGAGGT CCAGGAACTA AAAAGTCTGA TGTCCAATCC TGAAAAGCTC
27541   GATGGTGCAC TAGAGGAGGC TTTTACATGT AAGAGCATCT AAGTTCTGGA AATGCCAGTG
27601   TCAGGGAAGG GAAGTGGAGA GCAATTTGGC ATCCAAACAT AACTTGCTGA TACTTTTTTT
27661   TTTTTAACA CAAGTACTAC ATTCTAGTCT TTCTGTGGTG TCATTGTAAC TATTGTTTCT
27721   TAATATGCTA TCCACTGACT TCAAGGGATC AATAAATAGG AATCAAGGTG TCCCAGAATA
27781   TGGATTAGGG GAGTTTTTTT TTTGTTGTTG TTGTTGTTGT TTTCATCTAT TCATTATCCT
27841   GTAGCTGAAA TTTAGAATTT TCTTCCATTG TGTGTGACTG ATAGAAATAA CAAATTTGTA
27901   GGTTATAGTT GTTGCAAGAA TCTGGAAATC GTGCTTGCTT ATTTCCGAAG TACTATTAGG
27961   TATATCAACA AAAACACACA TATTACGGTC AAGTGGTTTG ATAATTATTT TAATATTATT
28021   GGTCTAATAC AATTGTAACC CTATGAATTA CTTTAAGTAT CTTATTTATG AAAAGAATCT
28081   GTAAGTTTCA TCAAACTACC AGAGCATACC GAAGACTGAA AAATTTTAAG AATCCAAACC
28141   TTAATGGAAA TGTTGGAGGC TGCCCAATTA GGTTCTGAAT TCCACCTTCC TGAATCACAA
28201   ACTTGTTTTA ACTCTCAGTC TGAGGTAAAC TACGTTTCTC TTTAAACAGA CATAGTTTAA
28261   TTTTCCTTTG ATTTTTGATT TAGTATTCTT ACTGATCATC ATAAATAACC AATGCTAATG
28321   TTAGTCTACT TTGGACCATG GTATTCGAG AAACTTTGAA CAAAGTCCCC TGCAAAACTA
28381   TGCATTGCAT TATTTCACAT ACATTATGT TTTCCAGACG GTTCAATAGT ACCTCACTTT
28441   TCTGAACTTA TTTGTATAGT TTGGCATCTT TTTAAAAATT GTGTCCTATA ATGAAAGGTT
28501   GTAAACATTA TGTTTTAAAT TTGTATAGAT AAAATCAACC ACAGACCTTT CCTTGCTTGG
28561   ATGTAATTGC CATTGTTTCC CAATGAGTTC GGAATTACTA GGATTGTGCA AAAATATGCC
28621   TCACTTGCCT GACATAGCAG AGAGCCATTT TGCCTAAATG CTGTGCCCAG CAATGGACTG
28681   TCACCAGATT CTCATCACAT ACAGTGAGGA TGAACAACTA GCCTCTCCCA GCAGCTGGCC
28741   GGTCTCTCAA TAATATGGGA CTCCCTCAAG ATGGCTTCCT GCACCTTTGC TCCTCTAGCC
28801   TTGTATGTAT ACAAGGCTAG CATGCCTGGC ATACATAAGG TTAAAAACAA AATCAATAAG
28861   TTATGGTTCT TCCTCCAGTT CTGGGGATTA TTAGACCACT TTTTTGTTTT GTTTTGTTTT
28921   GGATGGAGCC TCGCTCTGTC ACCCAGGCTA GAGTGCAGTG GCACAATCTC GGTTCACTGC
28981   AACCTCTGCC TCCTGGGTTC AAGCAGTTCT CTGGCTCAGC CTCCCACGTA GCTGGGATTA
29041   CAGGTGCCCG CCACCACGCC CAGCTAATTT TTGTATTTTT AGTAGACGGG GTTTCACCAT
29101   CTTGGCCAGG CTGGTCTTGA ACGCCAGACC TCGTGATCCA CCCACCTTGG CCTACCAAAC
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29161   TGCTGGGAAT ACAGGCGTGA GCCACCGCGC CCGGACTTAG ACCACTTTGT TTTGGCCAAT
29221   AGGACAACAG CCATAGAACC CTCCGCAAAT GAGAGCTTGT CCCTAAAGAT GCTTTATTTA
29281   CATAGCTGTG TGCCGCATGA GCCAAAAGGT GATAACCTTT GTTCAACACG CGCCTCCAGC
29341   CCTTCGGTTA AGTCCAAAGT ACCATTCTTA GAATGCTCTA AAATACATAA TTTTTTTTTT
29401   TTTTTTTTTT TTTTGAGGA GTCTCTCTCT GTCTCCCAGG CTGGAGGGGA GTGGCGCGAT
29461   CTCGGCTCAC TGCAATCTCT GCTTCCGGGC TAGCTGGGCC TACAGGTGCA GACCACCACG
29521   CCCGGCTAAG TTTTGTATTT TTTTTGGTAG AGGGGGTTTC ACCATTTTGG CCAGGCTGGT
29581   CTCGGATTCT TGATCTCAAG TGATACACTA GCTTTGGCCT CCCAAAGTGC TGGGATTACA
29641   GTCGTGAGCC ACTGCGCCCA GCAAAATGCT TTTTGTGGAG CCAATCACTT TATTAGCGCT
29701   TACCTCTCTA TGCCTACTTT ATGCTTTGAA ATTTTGTCAC AGTGGGGCCG GTCATGGCAA
29761   ACACAATTCA TTCTTATGCA GGCTGTCACG GTTATTTCTG TCATCCAAAC TCATTCTCGC
29821   AACGCATTTC AGCTCTTTAA ACGACTTTGT GAGCGGCCCT GAAAAGGGCC TTTGGGTTTT
29881   TTTGTTTTTG TTTTTTGAAG TTCTCAGGAG ACCGCGTATT CTTAGATTCA GCCGCCGAAG
29941   CCATACAGAG TGCGCCCCTG ACGTTTCAGG GCATATACTA CATCCATGGC TGTGACAGTT
30001   TTGCGCTTGG CGTGCTCCGT ATAGGTGACG GCGTCTCGAA TAACGTTCTC TAAGAAAACC
30061   TTAAGCACAC CTCGAGTCTC CTCATAGATA AGACCGGAAA TGCGCTTGAC GCCACCGCGC
30121   CGAGCCAAAC GGCGGATAGC CGGTTTTGTA ATGCCCTGGA TGTTATCCCG GAGCACCTTA
30181   CGATGGCGCT TAGCACCACC CTTCCCCAAG CCTTTTCCGC CTTTGCCGCG ACCAGACATG
30241   ATTCCTATCG CAGTGGAAGG TATGAACTGA AACAGTTCCT TAAATACAAA CTTGGCGGAC
30301   CTGATTGAAA ACAACATGAG TTGGCGCGGT TTTTTTTTTT TTTCAAATTT GGTCACCGAG
30361   TGGGTGGAGC AAGAAAAACT GTTTCATTAT GGTTCATTGT TTTGATTGGC CAGTGACAGC
30421   TTGCTCTTTG TGGGAGTGGA AGGGTGTTTG CAAGTTGAAT GCGCTGTATT CCTGTCAGCT
30481   TAATGACGCT AAGCATAGCC CCATTCCACA TTTCTTTTTA TTTCCACTTG CTAACTAATA
30541   AATTACGGAA TAGTTTATTG GGGAACATAC AAATAATGTT TAAAGGAGGT CAGATTTATA
30601   GGTCAAGGGA TTTACCCTCC CAATCATTTT AATATTTTTA TTTAAACCAG GCATTTTGAT
30661   GGCCTTCTCT GTGCTGGACA AGGTATAAGT TTGGCTATGA AGTTTCACTC CTAAAGACCC
30721   TATGTTTTGG GAAGGCAAAA AGGTAGCCAA ATAATTGCAA ATTAAAACCT CATAAGTGCA
30781   AACTTCTTCC TCGTCACTTT CCCTATCTCG ATTCAAATAT TTGTTGAATG ACTCATTTTT
30841   CTGCAAAAGT CTGAGAGAGA CAGGGAATAT AAACTTAAGT CTGGATAATA TGTTTTCCCG
30901   GGACGCTCTT CCTGGTCTGC TGTGCCTGTT TGCTGTGCCT GAAATTCCAA ACACTCTTCC
30961   CTTCCCTCCG TTTTTAATCC CCTTTCAACT TGCTACAGCT TTAGAGAAAA GAACATACGT
31021   TTTGTACAGT TGGGGATTAA TTGAAGTGTA GGGCTAATAC TTGATTAAGG TCATTACAAA
31081   ATCTACAGGG TCTTCCTCTG GGAGGTTTTT GTGATAAGAT TATTGGTGTT AAAATAAGGC
31141   TAATCCCCTT GAAAAATAAA TAGAATAGCA GAATTGGGTC TGAATGTGGT TTGAAGAAAG
31201   GGACTTCTCA ATTCAAAATT TTATTCTTAG CTTCCTGTGG GAGCTTTCCA GAATGCCCAT
31261   AAGATCCACT TTTGTTTAAA AAACAAAAAC AACCCCACCC ACCACTCTCT GGTTAATAAA
31321   TGAATTTCTA TTGGGAATAT TTAGAATGGG GCTGTGGCCT GTGAGAGACA TTATATAGTA
31381   ACCTCAGACT TGCTCACATG AAGAGAAGAA ATCCAGGAAT GGAGAAAAAA GACCCAGGAA
31441   AGGCCAGAAT GCTCTACATG TCATATTGTT TGTATCACTT CTGAAATAAT TGATTACATT
31501   CTTCTGCCCC AAATTGAGTT CTTAGGTTCT TCCACTCACT GTCCACATGC CACAACACAG
31561   ACCTTATAAC TAGAGACTTA GCTAGGAAGA AATGTCAAAC ATTACAGAGA AAAAATGCAG
31621   AGTCTGAGAT CATAAGTAAA ACTCTGAAAT CTCAACATGC CTTTTAATTC ATGAAAATAA
31681   AAAATATAGC AGCATATGCA ATATGATAAT TCTCTGAAAA CATACATCAT GTGAACTACC
31741   CTGGAACACA TCTCGCCAAG TGCCATCTTC ATTTAACCA GAGGTCTAGG ATGCCTTTCC
31801   TTTATTTTGC CTATTATATC ATTTATAAAA CCCCATTTTT ATTTTGATAT TTTATTTACT
31861   TTCTATTTCC TGCTCCTAAT ATCTCCTTTC TAAACTTTTC TCAATGACAG TGACTCAAAA
31921   ACAATGAATG TCAGAACAAA TATTTAAAGG ATCTGTACAT GTAGATATAT ATATTTAAAA
31981   TGGATTCTTC CACTCTGGGA AGAATTCAGG CATACTCAAT CTTATGGTTA GGGAGAGATT
32041   AGGCTCACTC GCCTAATCTG TATGGCTTCT CGTTCGCTTT CCATTTCACC TTCCTCTCAC
32101   CCATCAGATC AAACTCATTC ATTGAACAAG AGACCTAAGC CCTTCAGATT AAAACTCTGC
32161   AAACAAGTTG TGGTTGAGAG GATACATGAA GCATTCAAAC AAATAAATCT ATGATATTAA
32221   TCAGAGGTTA ATCTATGATA TTAATCAGAG GTTAATGCAG TGGCTCACGG CTGTAATCCC
32281   AGCACTTCAG GAGGCTGAGT TGGGAGAATC GCTTGAGCTC AGGAGTTCAA GACCATTTTG
32341   GGCAACATAG CAAGTCTTCA TCTCTACTTA AAAAAAAATA ACCAGAGGTG TTATGAAAAT
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32401   ATAAATTGTC CAGAACTACC CTCCACAAAC TAACTCTCTC AGAATATTCG ATATGAGGAA
32461   TGAAATATGG TGTGTGTGTG TGTGTGTGTG TATGTGTGTG TGTGTGTGTG TGTATGCACC
32521   TATATATGGC ACCTATATAT TCAACAAACA ATTCTGATAA TTGGCCAGGG TTGAGAATGA
32581   CTAGCAGCCC AGCATACACT ATCAGTTTTA AGTATATAAT TGCGCTTTAG TAAAATGTAA
32641   AGAAATCCCA GAGTAGAAAT ACTTTTAAGC TATATTACAG GTGAGAAAAT GCATAAGTAT
32701   AGTCTCACCC AACTTAGACT ATGGGGGCTT TATAATGTCA CAACAGTTGT TTCCAGGCAT
32761   TTGGGGACAT CACCACTGGT CTTGGGCAAG AAACTCCTCT AGCCAATGGC TGATTTATCT
32821   CACTCCCATC TAAGGCTTCA CTGCATTTCT CTTTTTCAGC AACCTAACTT ATTTAAAAAT
32881   ATCCATTTTC TGATTCATTT TTTTCTGAAT TAAACTGTCA GTACCATTGG CACACCTTTG
32941   GTTCCGTAGC ATACCTGTGT CTCTGCTGTG GTTTTTTTTA CCTCCACTCC TTACTTTTCT
33001   AGAAAAAAAT CTCTGCTTTT TCTTTTCAGT TTAAATTATT TCACAAAAAG TTTTCTTGAC
33061   TTGCACTTCC TAGGCTTGCT GTCCTTGTGT GGGCACGCTC CCATAAACAC TATTAATACA
33121   CTTCGATTTG TTAAAAATAA AGATATCTGG ACAGAAAATT TCTTTTCTTT TTTTAAGATT
33181   TTAAAATTTT TAATGTTTAT TTTTTTCCTA GACTGGAGTA CAGTGGCACC ATGATGGCTC
33241   ATGGTAGCCT ACACTTCCCC GGGCTCAAGT GATCCTCCCA CCTCAGCCTC CCAAGTAGCT
33301   GGGACTACAG GTGTGCACAA CCACACCTGA CTAATTTTGT TTATTTGTTT GTTTTGTTTT
33361   TTGAGATGGA GTTTCGCTCT TGTTGCCCAG GCTGGAGTGC AATGGCGGGA TCTCGGCTCA
33421   CCGCAACCTC TACCTCCCAG GTTCAAGCAA TTCTCCTGCC TCAGCCTCCC GAGTAGCTGG
33481   GATTACAGGC ATGCATCACC ACGCCCAGCT AATTTTGTAT TTTTAGTAGA GACGGGGTTT
33541   CTCCATGTTG AGGCTGGTCT GGAACTCCTG ACCTCAGGTG ATCTGCCCGC CTCGGCCTCC
33601   CAAAGTGCTG GGATTACAGG CGTGAGCCAC CACGCTCGGC CACTAATTTT GTATATTTTG
33661   TAGAGATGGG CTTTCCCTGT GTTGTCCAGG CTGGTCTTGA ATTCCTGGGC TTAAGTGATC
33721   TGCCCACCTT GTCCTCCCAA AATGCTAGGA TTACTGGCGT GAGCCACCAG GTCTGGCTGG
33781   AAAGATAATT TCTAACATTA TCCTCTCTTA AACATTTGTT TCAAAAATTT TACAAACATG
33841   AGAGTAATTA AATTTGATTT TCAAAATTCC CTTGAATACT TTCTTAATAG CACACAGAAA
33901   GCACAAAGTA TTTTACATTT GTTTTAATGA TGAAATTGTG AACCCAAACT TACACAAAGA
33961   AAAACCGTAA CATTATACCC ATACTTAAAA CAGATGCCCT CATATACATA GTAAAACTCT
34021   TGGGGGCAGT AGTGAAGTTG GTTATTTACT GTTTTATGAA AGTGCCATTC AGCCGGGTGC
34081   AGTGGCTCAT GACTGTAATC CCAGCACTTT GGGAGGTCGA GGCAGGCTGA TCACGAGGTC
34141   AGGAGTTCAA GACCAGCCTG ACCAAAATGA TGAAACCCTG TCTCTACTAA AAATACAAAC
34201   ATTAGCTGGG CGTGGTGGTG TGTGCCTGTA GTCCCAGCTA CTCAGGAGGC TGGGGCAGGA
34261   GAATCGCTTG AACCTGGGAG GCGGAGATTG CAGTGAGCCG AGATCGCACC ACCGCACTCC
34321   AGCCTGGGAG ACAGGGCGAG CTCCGTCTCG AAAAAAAAAA ACAAAAAGT GCCGTCATAG
34381   TGACTTAGTT TTAAGGAATA AATCAAGGAT ATTTAACTCA ATAGACTACA GTTAGCTAAC
34441   GTGACTTGCA CTGAAAGTTA TACGAATATT GGTACTTATT CCCCTGCCCC TGAAGTATGA
34501   ATTAAAGACT CCAAAATTCT TTTTAGAATC TTCAGAGTAA AAGCTAGAAT TTGATTTTTT
34561   TAAATAATAA AAAAATACTT TGTATCTAAA TCTGGTGTAT AAAATAACTT GGTGGATGAT
34621   GCTTCAAGGC TATCCATCCC CAAATTTCTC CCTGAATGAT AAAGAGAATA AATGAATATG
34681   TCAATTCAAA AGTTAGAAAT TTGGCCGGGC ACGGTGGCTC ACTCCTGATA ATCCTTTCGG
34741   ACGCTGAGGT GGGTGGATCG CATGAGCTCC GGAGTTCAAG ACCAACCTGG GCAACATAGC
34801   CAGAACCCGT TTCAATAAAT AATAGAAAAA AATGAGCCAG GCGTGGTGGT CCCAGCTACT
34861   CAGTAGGCTG AGGTGGGAGG ATCACTTGAG CTCAGGAGGT CGAGACTGCA GTGAGCCGTG
34921   ATCGCAGTAC TGCACACCAG CCTTGGTGTC AGACTGAGAC CCTGTCTCAA CAACAACAAA
34981   ACAAGTTAGA AATTTGGCTG GGCGCGGTAG CTCACGCCTG TAATCCCAGC ACTTTGGGAG
35041   GCCAAAAAGG GCGGATCATT TGAGGTCAGG AGTTCGAGAC CAGCCTGGCC AACATGGTGA
35101   AACTCCATCT CTACTAAAAA TACAAAAAAA CTTAGCCGTG CATGGTGGCA TGCGCCTGTA
35161   GTCTCAGCCA CTTGGGAGGC TGAGGCAGGA AAATTGCTTG AACCCAGGAG GCAGAGGTTG
35221   CAGTGAGCCG AGATCATGCC ACTGCATTCC AGCCTGGGTG ATAGAGTGAG ACTCCATCTC
35281   GAGAAAAAAA AAAAAATTCT GTATGAACTG AACAAAATAT CCTTAAATTT TAAAATACAT
35341   CTGAAAGATA TTTCAAAATA TTTAGGAAAA AAATTATAGG GATCAGGCAA ATTCTGAGAT
35401   TCCTTTTTCC CTGCAGCAAA CATTAGGAGT GCTGCTGTTC CTAAAAACAT GGTAACTGTT
35461   GCCACACCGT ATGTTCCTT GGCTCAGACA TAAGGTTGTG TAGTTGTTAT CCAGAATAG
35521   CTAGAATAAA AATCCAGCAC ATCATTTTCT TCAGCAAGTT AACTAACCTC TCTGTGCCTT
35581   GGTTTCATAA CAGCAACATA AGCATAACAG AATAGCAGCA ATAGCTCCTA CCTACCTCAT
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35641   AAGATTCTTT GGAAGAATTA AATTAAGATT CAGAACACAG CCTAATATCT AGTAAGTAAT
35701   AATAATTGGC TAAAAAAATT TTCTTAAGAT TATATATATT CATGGGGTAC AAGTACAATT
35761   TTGCTACATT AATATATTGC ATTGTGGTGA AATCAGGGCC TTCAATCCAT CCCGGAAAAA
35821   AAAAGTTTTT GAAAAGATTT CTGCCATGGA AAACTTTTAA TGTACAAATT CATCCATCCA
35881   AGAAATAGAA AATATATAAG TATCAACTCC AAATCCACCA TATCTATCTC TTCTGCACCT
35941   TAAACAATTA CTCAGAAATA GAATGCTTGA GATACCAGAA TGCATGCATA TCAAGTAATA
36001   AATGCATGCA GGATGTCAAC GCATCCTAGG CTTTCAAATA AAATTGTCAT ACAAAATACT
36061   TTAATATTGT AGTAACATTC TACATGTTAG AGTGTAGAAG TTAATCGCTG ATGCAAAAAA
36121   GGAAAAGAAC ACATTATACC CAAAGCCTAC AGAGAGAATC ACAATTACAA ATATCAGCCT
36181   GCATGTGAAA ATCTTTAATT TGAAAGTCAG AAATATTTAA ATGATAGTCA TTGTTAAATC
36241   AGATTGTGGT TTGAAAAAAA GTTAGTTTAA AACTGAGTTT ATGAAAAATT TGGGGATTTT
36301   AGAGACAGTG TTTTGTTTTT AAATGTGTGT GAGTTTGTGA AGAATGTTTT ATAAAATACT
36361   GACAGTATTA TAAGATGACA TTATTATAAT ACAACATAAG AATTTTGGCC TGTACCTCTC
36421   AGCAGTCCTC AATCACCTGC TGTACTTGAC TCAATGATTA TCAGAGTGGT TTGTTTTCCT
36481   TCTGTTGTGT TCCCAGTTCA GGCAGCTCAG CAATGGCCTG TGATTCCAGC AATTCAAATA
36541   GCTGGTAAGT AGTTCTTGT TTGTTTTCTC AAATTTTCAG GGGCTTTTCT CTACAAGTGA
36601   TTTCCAGTGC ACGCCCTCC ACCCATTCTT TATTCCTTTA CCTTCAGGAA AACCCTCAGC
36661   GCTGCATCTC TGGTCACCGG ACCACCGTGG TACATTTACC TATGGCCACC AGGTGTCACC
36721   CTTCTCTTTA CTACCATGGT TTGTGAATGG TTTTGCCAGA GGTGAATAAG AATTTAAAAT
36781   GCAGGTCTTT GATTTTTCAA ATGTAGTTGA CCTTAAGAAT TTATGAATAA AGCCAGAAAA
36841   ATTAAGCTTA AAAAACACCG AAAGAAAATG AGGACTTAAA ATTTCTATTA AAAAAATTAA
36901   CAGGCCACAG TTGCTGATGT TTAGTAAATG TGTTAGTGAA ATGTGTTACT GTGAAGACTG
36961   GGGTGTTTCT TGAAATCTCA GCCCAGGTGA AATAAAACCA ATATAAAACA AATGCTTACC
37021   TAATAAATTA ATTGTAACAT ATTCCTTATG AGGTAGAAGA GTAAGTGAAG CCTTATAGCA
37081   GTCTGCTTTC AGTATAGTAA GATATTAAGA GAGAAATAAT TTGTCATATG CTTTCAGAAT
37141   GGTTTGCTGG TAAAATAACC AATGTCTTAC AACTTAGACG ACAATGTCCC TAGAGTGAAG
37201   AAACACGATT AATTCGGCTA CCACAGTTGA ATGAAAATAT TCCGTAAGAC AAAATGTAAA
37261   GAAATTAGAA GCAAAATAAA TGTCTCCAAA ATGACAAAGC GATTAAGTAT ATACACAAGA
37321   TGAACAAGAA CTTCAATAAA ATCATGCAGT ATACAATACA ATGTACATTT ATTAAAGTAT
37381   ATGCATTTTT AATGCAACAA TAATACTAAC AGGTAATAGA CAAGTTGTTA ATAGTTTTTC
37441   ACTGGCTAAT TAAATAACAG CTTTAATTGT ATTCATTTTA TAGCTTTTCT ACAATGAGCG
37501   TAAATCACAT TTACTTTTTT CTACATAACT TTTCTAACCA CAAAAAAAGA AAATGGTTTA
37561   AAAGAAGAGA TGAGATATCT TTGCTAAAAT TTAATGCCTA AGAAGAAAC TTCTGAGCTG
37621   TATATGGTAT CCTGAAGCAC CTGCCCTTCA AGACAGAATG CTTGTACCAC ATTTATGCAG
37681   CCAAGTGCAT GTAGTAACAT AAAGTAAACA CATGCCATCT GGATATATAT ATTAAGACTC
37741   TTTTGACGGC TGGGCAGGGT GGCTCACACC TGTAATCTCA GCACTTTGGG AGGCCGAGGC
37801   AGGCGGATCA CGAGGTCAGG AGAGTTCGAG ACCAGCCTGG CCAACATGGT GAAACCCTGT
37861   CTCTACTAAA AATACAAAAA TTAGCCGGGC ATGGTGGTGC ACGCCTGTAA TCCCAGCTAC
37921   TTGGGAGGCT GAGACAGGAG AATCGCTTGA ACCTGGGAGG CAGAGGTTAC AGTGAGCCGA
37981   GATCATGCCA TTGCACTCCA GCCTGGGCAA TAGAGTCTCA AAAAAAAAAA AAAGACTCTT
38041   TTGAACATGG TGAACTGATT TCCCAGAATC TAGCAATTCC TGAATGTCCT GGTTAGATTT
38101   TTTTTTTAAT GTGCACCGGA ACCCCAGTGG CTCCATGGAA GGACCTGGGC ATCCTCTAAG
38161   CCACTTGGTG GCTTCCATTA TACCATCTCA AAATGAGAGA GCTTACTCCA CTTCATTGAG
38221   GGAAATACCA CCAGAGTTCT GACTCCAGAG GCACTGGCCT AGGGAGGACA CCGTGTGTGA
38281   AGCCCAGCAG GGCCACTAGC TGTCCCCACC AATTACAGTC CTTGCGTAGG GTCCAAAGAA
38341   ATGAATGCCA AAGAGAGCAA CAGAGGAGCA AGGGAGTCAC ATTCCAGGAC CTTCCTTCAG
38401   GGACTTTTAA AGGAAACATG ACAGCTGAGG ATCAGTTGGT TGTTTTCTGC TGTTCCCCTT
38461   CATGTGATTC AAGCTCATTC AGAAGAAACA CAATGAGACA AGAGAAGAGC CATCTCCTTC
38521   CTTCTCTATT TATTCTAGGC ATCTAAACTA CTGAATGTAG TGGTGTCTGA GATGTATCAA
38581   ACGGTCAGAT TGACTGAGTT TGAAACCTGT TTCTATCACT GACAAACTAT GAGATACTCT
38641   ATACTTCACT TTCTTTTTTT TTTCATTTTT TTATTTTTAT TTTTATTTTT TTGAGATGGA
38701   GTCTCACTCT GTCACCTAGG CTGGAGTGCA GTGGCGCAAA CTCGGCTCAC TGCAAGCTCT
38761   GCCTCCTGGG TTCATGCCAT TCTCCTGCCT CAGCCTTCCG AGTAGCTGGG ACTACAGGCG
38821   TCTGCCACCA CGCCCAGCTA ATTTTTGTA TTTTTATTAG AGATGGGGTT TCACCATGTT
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38881  AGCCAGGATG GTCTCGATCT CCTGACCTCG TGATCCACCC GCTTTGGCCT CCCAAAGTGC
38941  TGGGATTACA GGCGTGAGCC ACCGTGCCCG GCCTACTTCA CTTTCTTCAT TTAAAAAAGA
39001  AATGGGGATA ATAGTACCTA TCTCATAGAA TTATTGTAAG AAGTGCATGC AGTAATGCAT
39061  GTAAGTAGGT GCTCAGAAGA GTCGGACACG AAGTAAGTGC TTTTATCATC CTTATCATAA
39121  TTTTCATTAT CAGAACAAGG AGAGACCAGG TAGAAAATTA TTGTGATTCT TCAGGTCTGG
39181  AATACTAGAG TAGCATCCCA AATGAAGGCA CCATTAAACT TTGCAAATCT GTATGACACC
39241  TTCATGCCAA TTAGAAAAAA CACCTCTTCA CAACCCCTTT CAAGATATTT GCCTCCTACC
39301  TGCTAAAAAC ACCCATCATA CTACCCACAG ATAGCCATGA TGCTTTTTCT GGGACAGGTG
39361  CCTCTTCCAT TCGTGCAGTG TACAGCCTTC ATAGCTGTGC AACTCACATC ACAATCAGAT
39421  GGAAGAATCC CCAAGGCTTG GTGACAGATG AGTTACTGGG TAACACAGAG AGAGGATTCA
39481  AAGGAAAAGT TGAACGGGTC CAGAAAATGC ATAGATACAT GTGTAAAAAT CTGGTAAGGT
39541  TATGACTAGC CACGTCCCAG GGTTCAAAGC TTTTCTCAGA TGTTAAAATG AATCATGTAA
39601  GTCCCCCAAA TTTAAGGAGT CCTCTTCCAA AAATAGGAAA TGAAATGACA TAGGTGTATG
39661  TCTCTGAGGT GACGGAGGAA ATGAAGGAAG CCTCTAGATG CAGCTTGAGG TTCATGAGAG
39721  ACAGTTCCAG GGGAGAGGTC ACAGCTAGGG ATCACCGGCA TGCAGGAACT CAGAAACCTA
39781  AATGGGGAAA TCTTTTTGAG GAAATGAACA GAGAAGGCTA AAATCAAGGA GTTCGTCAGG
39841  CAATTTCTAT GTTTAGGTTC AACTCTCTCC TGAAACATGA AGAGCTCATA AATGCACTCC
39901  CTCTTTGAGT CTCTAGTTTT GTCTCCTTCC CACAGTGAGT CTGCAGGCTG CGTGTCACTC
39961  ACGTTCAGCT AAGACGTAGT GCCCCATGGC TCCTCCTGTG GAGACAAGAG ACCCAGGAAA
40021  GAGGCATCAC AAACCTAGGC ACCATCTTGC CTCTTCTCTC TTCCTTATTT TCCTCATTCA
40081  CCCATCTCAA TTTAGACCTG GGCACTATTG GATTTCAAGA ACCATTATCT CTCATCTGGA
40141  AATGCTTATT GGCTTTCTAA CTGGTCTCCT CACCTCTCAT CTAACTTCTT AACAACACAT
40201  TCACCATATA AGGGAGATCG TGGTCCTCCT TTCTTAGGAT CCTTCAATGA CACCCCAGTG
40261  ATCATAACCC AATATCCCAA AAGACCCTTG GACTCTGTAT GAGCTGGCTT CTTTCTGATT
40321  CTCTTTTCCC TACACCACAG ATGTTCAGGG GGTAGAAATG CATAATTGGT GAGTGATAGC
40381  TAAGCAAACT CAGGGTTAAG GTACAGTAAT TATTTCTAAT CTCCCAGTAT GCCTTATACT
40441  CTCCTACTTG GCATGGTTGC TCCGTCTGTG TAGACCTCCC ATCATCTTCA ACCTCACCTA
40501  ATGGAATCCA GCTTCTCCTT CAAGATCCAG AAGGCTATCT TGATCCCCAG CTGAATGTGA
40561  TCATTCTTTC CTTTGACACC CTAAGCATTT GCTTCCTGCC TGCTTTAGGA CCTCATGGGG
40621  TCTTCTTTAA CTACATTTAC TTGCTATCAA TTTCATTCCC TACCAGATTT GGGTTCTGAG
40681  AATAGCCACA GTGACTTCTC AACCTCAAAG CCCCTGTACT ACCTTAAACA GCTCTTGCAA
40741  AATAGTAGGT GCTCTGAAGA TGTTTGTTGA ATTAGAGACT TTCATTCTGG GGAGAACCAT
40801  TATTTTCTGT CTCCCAGGGA GCTGCTGGTG TCCCCAAAGA ATATAAATGA GAAAAATGCT
40861  TCCCATGGAT GCCAGATCCC CTCTGCCCCT CTTCCCACTG TGCCCTGGGG CAGAGGTACT
40921  AAGAGACTTC CCCCTTGTTC CTACTCACTT GAACCCTGCC TCTTCCTTAA TATTATGAAC
40981  AAAATTCCAA TGAACAAGAT GACGACAAAA ACAGCAATTC CACTGATGAC TCCAATGACT
41041  AGGGTGCCAG ACGGTGAGGG CTCTAAAACA GAAAAAGCAA GTTAAAGCCT TTGATTGCCA
41101  CCCTCAGCCC ACCCCCTAAC AAAGAGCAGA TCCTCATCTC ACTGCCATAA TTACCTCCTC
41161  AGGCACTCCT CTCAACCCCC AATAGATTTT CTCAGCTCCT GGCTCTCATC AGTCACATAC
41221  CCCAGATCAC AATGAGGGGC TGATCCAGGC CTGGGTGCTC CACCTGGTAC GTATATCTCT
41281  GCTCTTCCCC AGGGGGTACA GCCAAGGTTA TCCAGCCCTG GTAGGTCCCA TCCCCATTGG
41341  GCAATACGTC TTTAGGTTCG AACTCCTTGG CATCCATTGG CTGCTTATCC TTCAGCCACT
41401  TCATGGTGAT GTTCTGGGGG TAGTAGTTCA AGGCCCGACA CCGTAGAGTG GTCACTGAAG
41461  AGGTCACATG ATGTGTCACC TTCACCAAAG GAGGCACTTG ACAGGAAAGA GGAAGGATGA
41521  GGAGAGGGGA TCTGTTTACC CTTGCCAGGA AGACTGGAAC TTTCACTTCC TTCTATAGGT
41581  TGGAGGAAGG AAATACCCTT TTCAGAAAAA AACAAGCTAC AGGAGAGACA CCATTTTGTG
41641  TCCTAAGATT GGACTCTAAC ACAGTGTCAC TTGGAGAGCA GTCAGATCAG CTTGTTCTCC
41701  TCACATGTAA ATATACATAT CTGTTACCCA TGTTCTTTGT TCTGATAGAT AAAATTGCCC
41761  TTTATGTGCA TTGAAAATGA TTGAATACAG ATGGTCAGTT TCACCTGGGT CAACCTAGGA
41821  GGCATTGTTA TAAGAAGCGG ACTTGTAAGA TAGGTAGCTT CAGTGATTAT TGCTATGTTC
41881  TATGAAAGAA ACTTTTAACC TAAAGGATTC TTCTACTCTG ATAAGTGGCC TCACTTGATA
41941  TTTTGTCCTG GTATTCATAT GATAGCTGAG ATCTCTGAAT TCTCTTTTTT TTTTTTTTT
42001  TTTTTAAGAT GGAGTCTCAC TCTGCTGCCT AGGCTGGAGT GCAGTGGCGC GATCTTGGCT
42061  CAGTGCAACT TCCGCTTCCC AGGTTCAAGC GATGCTCCTG CCTCAGCCTT CCAATTAGCT
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42121   GGGACTACAG GTGCGCATGA CTGTGACCAG CTAATTTTTG TATTTTTTTA GAGACGGGTT
42181   TCACCATGTT GGTCAGGCTG GTCTCAAACT CCTGACCTTG TGACCACCCG CCTCGGCCTC
42241   CCAAAGTGCT GGGATTACAG GGGTGAGCCA CCGTGCCCGG CCTTGACATT TCTGAATTTT
42301   TAACAGGTAT AAATATACAA AAGATTATTG GTTAAATAAA AAGCAAGGGC CATAGACACT
42361   TCCCTTTGAG CCATATGCAT GGAGAAAAGA AATTAAACCC ATGACTTGTG GCTGTCTCAT
42421   ACATCTCAAT TATAAGGTAG AGACTCTAGG ATTGAGAAAG TCCCTTCCCA GAATTTGGAG
42481   AGGCACACAG CCTCAGCCAC CTCTGAAACT CCAACCAGGG ATTCCGTGCC CTGCAACCTC
42541   CTCCACTCTG CCACTAGAGT ATAGGGGCAG AAGTGTGTTT CCACCATACC TTGTTGGTCC
42601   AAAACACCTC TCCCCAGCTC CAGCAACTGC TGCAGCTGTG CAGGGCAGTC CCTCTCCAGG
42661   TAGGCCCTGT TCTGCCTGGC CCGAATCTTG TGCCTTTCCC ACTCCAGCTT GGTGGGCCAG
42721   GCCCTGGGTT CTGCTGCTCT CCAATCCAGT GTGTCAGGGC AGAATTCAAG GTGGTCCTGC
42781   CCATCATACC CGTACTTCCA GTAGCCCTCG GTACTGTTGT CTTCTTGCAT TTCACAGCCC
42841   AGGATGACCT GCAGGGTGTG GGACTCTGGA AAAATCCCCA GCCTTGTTAA CTGCAACCAA
42901   AGGAATAGGT CCCTATTTCC ACCATCCCCA AGGACCAAAT GATCTCAGGA AGCAAATTCC
42961   TTCCCTCTTC CCTGCTCCCA CAAGACCTCA GACTTCCAGC TGTTTCCTTC AAGATGCATG
43021   AAAAGATGAA AAGCTCTGAC AACCTCAGGA AGGTGAGGCC CCCTCTCCAC ATACCCTTGC
43081   TGTGGTTGTG ATTTTCCATA ATAGTCCAGA AGTCAACAGT GAACATGTGA TCCCACCCTT
43141   TCAGACTCTG ACTCAGCTGC AGCCACATCT GGCTTGAAAT TCTACTGGAA ACCCATGGAG
43201   TTCGGGGCTC CACACGGCGA CTCTCATGAT CATAGAACAC GAACAGCTGG TCATCCACGT
43261   AGCCCAAAGC TTCAAACAAG GAAAGACCAA GGTCCTGCTC TGAGGCACCC ATGAAGAGGT
43321   AGTGCAGAGA GTGTGAACCT GGAGACAGAG CAACAGGCCT TAACCATGTG TAGTAGGAGG
43381   GGAGCAGGAT GTTGAGGCTC CACACACCTG CATCAACTCA TACCATCAGC TGTGTCTGGT
43441   CCTCATTTTG TGAAGGGTGA GTTGCAGTCC TGTCTTTCTT CCATATGACA GTCCTGGGTG
43501   CTCTTTCCTT GTGTGCTTTT CTCTGCCACA CGTGGCTGCC ACCCCCTCAC TGCCCCCAGA
43561   TCCTATTCCA ATACTCATGA TTAGACAGAC TCCACTAAAG CTGGTGGATT CTAGAAAATG
43621   TTAAGGTGTG TCTAGCCATG GTAGTTGAAC TCAGGAGTTG GTGCTCAGGG CAAATTAGAC
43681   CCAAATCCTG AGGAATAATT CCTTCAGTTT TTTTTTTTTT TTTTTTTTTT TTTTTTTTTT
43741   GAGACAGAGT CTCACTCTAT CACCCAGGCT GGAGTGCAGT GGCACAATCT CAGCTCACTG
43801   CAACCTGCAC CTCCTGGGTT CAAGGGATTC TCCTACCTAA GCCTCCTGAA AACCTGGGAC
43861   TATAGGCGTG CGCCACCACA CCAGGCTAAT TTTTGTATTT TTAGTAGACA TGGGGTTTCA
43921   CCATGTTGGC CAAGCTTGTC TCAAACTCCT GACCTCAAAT GATCTACCTG CCTCAGCCAC
43981   CAAAGTGCTG GGATTACAGA AGTGAGCCAC CGTGCCCAGC CTTGGTCCTG AATTCTTACA
44041   CTGAACTGCC TATGTGGCCT CACCACTTGG AAGCCTGACT GGAATCTCAA ACTTAACATG
44101   TCCAAATGCA GATCCTTGAT TTACCCCAAA CTGCTCTTTC CTCTGCCTTC ACCATCTCAG
44161   AAATGGCATT GCCAATTACC CCACTGCTCA GGCCAATAAA ATTAAAATAA AGAACAAAGT
44221   CAACTTTAAC TCTTCTCTTT TTCAGGGGGT CAGGGGAGAC AGGGTCTTGC TCTGTCACCT
44281   AGGCTGAAGT ACAGTGGCAC AGTCATGGCT CACTGCAGCC TCAACTTCCT GGGCTCAAGC
44341   AATACCCTCC ACCTCAGCCT CCCGAGTAGC TAGGATCACA GGTGCATGCC ACCACACCCA
44401   GCTAATTTTT GTATTTTTTG TAGAGAAGGG GTTTTGCTGT GTTGCCCAGG CTGGTCTTGA
44461   ACTCCTGAGC TCAGGAATCT GCTCTCCTTG GCCTCCTCCT TGGCATGAGC TACTACACCC
44521   AGCCAATTCT TCTCTTTCTC TCACACAACA TAGAATCCTT CAGCAACTTC CTTCAGAATA
44581   TATTCAGGAG ACAATGGTTT GTCACTCCCT TTTCTGTTCC CACCCAGCCC ACTCCACTAC
44641   CTCTTGCCTG GACTGTGTAA CAGCTTCCTG GCTGGGCTCC CTGCTTTTAC TGTTGCTCCC
44701   TTCATTCTGC TTTCCACATA GCAGCCAGAG CAATCTTTTA AAAGCCTGTG ACAGATCACT
44761   GTTACTCCTT GGCTAGAATT CACACCACAG CCTACAGGCG CCTGCACAAC CTTGTTTGTG
44821   GCTCCTCTTC TGAGCCCATT ACCTACTTCT GGCCTCTAC TCCCCAGCAC TACTTGTTTA
44881   TTTTTTTCAA CCCGAGCTTC TTAACCAGGA GTTTGTCTAC TAGGTGACAT GTGGCAAAGT
44941   TTAGAGACAT TTTTGGTTGT CAAGACTGGG GGAGTGCTCC TAGCACCTAG TGAGTAGGGA
45001   GGACAGGATA CTGCTAGACA TCCTACATGC AGATGGTAGT CCCCCTTCCC ACCCCCACGC
45061   CGCCCCCCCC CCCACACACA CACACATGAG TAGTGCTGAG AAAACCCGCT TTTTAATCCA
45121   ACTTGCCAGG CCCACTCAGT TTGCCTGGGA AATACTGCTC CCAGTCAATA TCATTCTTAT
45181   TTCCTTCATG TCTCTGCTCA AGTGTCAGCC CCAGAGTGAC TTGCCCTGAC TTCTCTGCTT
45241   CTCACAACAC CCATGATTTC CTGATGTTGT ATATCTTTCT GCTCATTTGC TTATTGTCAT
45301   CTCTCCCACT AGAATGCAAA ATATCAAAGG GTAAAGACTT GTTTCCCTGC TCTCTCCCTT
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45361    GGGGCTTGAA  CAGTGCAACA  CATGGCTGGG  ACTCATTTAC  ACTTGTAAAC  AATGAATATT
45421    TCTGCTCAAC  ATGAAATTTT  ATTATTCAAC  CTCTAATGCA  GTGTGATGTT  TAAGAATCAT
45481    AGCTATGAAG  TGGAGACATG  AGCTCTGCCA  CCAAAGCCCC  GTGTACCATT  GAATAAATTT
45541    GCCAGGAAGC  AGGCCGTGCC  ATGCCTCATT  CTTGTCATGT  GTAAAATGTG  GATACACGTA
45601    GTACCAAAAC  TCAAAGTGCT  GTGCTGAGGC  CGGCGTGTGA  CCCACAGAAC  ACTGTGCTAC
45661    ACTACAGGGC  AAAATCACTG  TCAACTAAGA  TTAGAAGCAG  CTGTAGTACT  TGAAATAACA
45721    TCAGAAAACC  AGATTATTTA  TGTTCTTTGT  AACCTGAAAA  GAGTTATATA  ATCTGAATTC
45781    CAGTTAACTT  CTAGTAAAAT  AAACGTATTA  TTAGCTCCTA  CCTCCCTATG  CCTAGTGAAA
45841    ATCAAATAAG  ATCAGATATG  AATGTAACTT  AGAAGTGAGT  GCATTGCTTA  CATGTTCATT
45901    ATCAGTACTT  TGTAGAGAGG  CCTCTTAATT  ACACAGCACA  TTGCAAATCA  ATAAAGCCTA
45961    GCCGAAAAGA  GAATTGTTCA  GTTCAAACGT  TCAAAACTAA  CATATACTTA  ATTTTCCAGG
46021    CAAAAGAACA  ATTGCCAAGA  GTGGGGAAAG  GCCCGAGGTA  GGCCTCTCTC  AGGAGCCTCC
46081    CACCCTAGAG  ACCTCCACCC  CAGGTCTCAC  CAAAAGTGGG  TGGAATGGTG  AAGAATTCAG
46141    ATCCCCAACG  CCACTCTTTC  GCGCCCCCAC  CGCCCAACGC  ATTCGTTCTG  AGGTGGAAAC
46201    CCCGTGCGGA  TCCTGCTGTG  GGTTTGCTCA  GCCTTCTCGG  CAAGCACTCA  GGGAAGAACT
46261    TCCTGTTTGG  AGATGACTGG  GGAAAAAACT  GCACAGCTGA  CATTGGAAAT  AAACCCGAGT
46321    TCCAGGTTCA  AGGAGCCCCA  GGCTTAGCTC  AGCTCAAGTG  AGGAACTACG  AGATTTATTT
46381    AAAAGCATTC  TAGTTGGGGG  AAGGGAGTGG  GCGGTTCCAA  AAGTCACTCC  GCAGAGCCGG
46441    GACAGCCGGG  GGAGGGGGCA  GGTCCTGGGG  CGAGGGACCC  CTATCTGCAG  TTCAGTGGTA
46501    GGCACTCCCT  CACGGGGTCT  GGACGCAGAA  AGTAGGGAGA  GGGGCTTGCG  GATTGGGTTG
46561    AGCAGGTCCT  CCAAAGTTAG  CAAACTCCCA  AGCGCAAAGA  AAAAGCTAGT  TTCGATTTTT
46621    CCACCCCCGC  CGCGCCCCTA  GTTCGCCCGC  AGCCCTCGGA  CTCACGCAGC  AAGCGCCCCT
46681    GCAGGACCGC  GGTCTGCAAA  AGCATCAGGA  GGAGAAGCGC  CGGCCTGGCT  CGCGGGCCCA
46741    TTTCCCCAGC  TCTGGCCGCA  CGTCCCCGTT  AAATCTCCGC  TTCTTTTGGG  GGGCGGGGAA
46801    ACGGGGATGG  CTCCAGAAGT  CACCCTACAG  CTATTGCCTA  GGCTCAGGAG  ATGCCCAGTA
46861    AAACTTCCTG  GTGAAAAGCA  ACAGGTCTTT  CAGAACTTTA  GTTCTCTCTC  TCCTACAGCA
46921    GAAGGTACCT  GCTTGTGAAA  CACTAGGTGA  TCCAGTGTCC  CCCTTGGTTT  TTAAATCCTG
46981    AAGGGGTGTT  GTTGATTGGG  GAAAGTAGCT  TCGCAATGTT  CTGATCTGAA  CTTTAGATAT
47041    TTAAATATTT  ATGATTTTCA  AAATTCAATC  ATACATTTAA  AAATTTATC  TCAACCTTAG
47101    ACCAACTTAT  GTCTTATTTG  ACTTAGAAAT  ATAAAGCTTT  TTCATTTTGT  TTTTTGATTC
47161    AAATTAATTA  AGTCATAACA  TTAACCAATT  AGATCCTACT  GAAACACGTT  CCACAGCCTT
47221    CATAATTGAA  TTATCTGACA  AGTGTTTCAC  AAACTTTACA  GTATTGGGAT  TATCTGGAGA
47281    ATGATTAAAC  ATATTGAGGC  CTGCTCCTAA  CCCCAGACAC  ACTGATTTAA  TGGGTAATTG
47341    TTAGGTAGTT  AGACATTAGC  AGTTGGGAGG  GGATGACAGA  AGAGAGCGGA  AAGGCTGTCA
47401    CTAAGACAGC  CACTGGCCCA  CCTAAATTCA  GGCCCAAGAC  TACCCTAATG  CCACCCTAAG
47461    GGATGGAGTT  TATGATAAAG  TCTGTGGCCA  AAATATCCTG  GAGAAAGAGA  AAGGAGGGTA
47521    CAGGTGGAAA  TTCCCTAAGG  TGGCACATGC  CCAACAACAC  AAAAGCCTGT  CTTCAAGTTC
47581    ACCCCAAGTT  CATCATGCCA  TCATTATAAT  AGAATTTACA  TACAGTTTTG  CCCCCCCATC
47641    CCTGGGAGGC  TTTTCTTAAC  AAATTATAGG  TAAGACCATG  CACAGTTTAA  TTTTAGATTG
47701    TATAGCTATA  AACTTCAATC  AAATAACATC  ATCCTGTCAC  TCAGATACAG  CCCAAACCTC
47761    AACTCCTCCC  CACAAACCCC  ATAAAGCAC  CTTGAGCTCT  GTAAAGAAGT  GCTGAGTTCA
47821    CTTCGCAGAA  ATAAGCCCGC  TGTCCCTCAG  AGTGTATTAT  TGTGCTTCAA  TAAACTTTGC
47881    TTTAAGCTTG  CATTTTGGTG  TTAGTTTGTA  GTTCTTTGCT  CACTATCACA  AGAACTGAGA
47941    TTGCTGCTTC  AGAGCTCCGG  CTATAATAAT  CTCCTCGGTT  AAAGGATCCA  TCCCAATGCA
48001    TAATTCCCAG  TAACAGTATG  GGATGCCACC  TGGGCAATGG  GATTTTAAAA  GCTTTCCTTC
48061    TCCCTCAACG  AAGTTTGGGA  ATTATTGCCT  TAGACATTTC  AAACAATATT  AATAAATTTA
48121    ATACACCTGA  TTTGCTCCAA  ACCTTTACAT  ATCTAGCAAA  TTCAACAGGC  ATTATTTTTG
48181    TAAGCATGTA  TGCAAATTTT  GGCAATTCAA  GAAAATCAAA  CAGGATATCA  GGGCCTCGAC
48241    TGTAGGCAAA  CAGATACAAT  AACATTGGAA  ACATGTAGAA  TATTGATGAT  GGGCACATTG
48301    GGGCTGATAG  TACTATTCCT  TTTTTTCAAT  TTTTGGTAAG  ATATAATTAG  CATACCATAT
48361    AATTCATCTA  TGTAAAATGC  AAAAATTGGC  CCAGCTCAGT  GGCTCACGCT  TGTAATCCCA
48421    GCACTTTGGG  CGGCCGAGGA  AGGCAGATCA  CCTGAGATCA  GGGGTTCGAG  ACCAGCCTGG
48481    CCAACATGGT  GAAACCCCGT  CTTTACTAAA  AATACAAAAA  TTAGCCGGGC  GTGATAGCAG
48541    GCAACTGTAA  TCCCAGCTAC  ATTAGAGGCT  GAGGCAGGAG  AATCGCTTGA  ACCCGGGAGG
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48601  CGGAGGTTGC AGTGAGCTAA GATCGTGCCA TCGCACTCCA GCATGGGAGA CAAGAGCAAG
48661  ACTTCATCTC AAAAAAAAAA AATTAGCTGG GTGTGGTGGC ATGCACCTGT AATTCCAGCT
48721  ACTCGGGAAG CTGAGACAGG AGAATCGCTT GAACCTGGGA GGCGGAGGTT GTGGTGAGCC
48781  GAGATCATGC CATTGCACTC CAGCCTGGGC AACAAGAGCG AAACTCCGTC TCAAAAATAA
48841  AATAAATAAA ATAAAATGCA AAAATTAATG GATTTTAGTA TATTTACAGA GATGTGCAAC
48901  CATTACCAAA ATTTTACATT TCTATCTCCC CAAAAAGAAA CCATGTTCCC CTAATTCAGT
48961  ACCCTTAATT CATCGCCTCC CAGATTCCTC CATTCTCCTC CTCCTCCCCT CCCAGCCCTA
49021  GACAATCTTT AATCTACTTT CTTTCTATTT GGAACATTTA GTATACATAG AGGCATATAA
49081  TATATTGCTT TGCCGTGACT GGCTTCTTTC ATTTAGCATA ATGTTTTTAT GTATGTTTTT
49141  CATGGACCAA TAATATCTAT TATAAGGACA TACCACAACA TATTTTATTT ATTCATTCAT
49201  CAGCCGATGG ACATTGGTTT GTTCTACTT TATGGCTATT GGGAATAGTG CTGTTATAAA
49261  CATTTATGTA CAAGTTTTTT TGTAGACTTA TGTTTTGATT TCTTTTGGTT ATATATCTAG
49321  AAGTGGGTTT GCTGGGTCAT ATGGTAACAC TGTTTAACCT TTTGAGGAAT TGCCACATTC
49381  TTTTCCAAAG TAAGCATTTT ATCCTCCTAT CAGCAGTGTA TGAGAGTTCT GATTTCTCTC
49441  CATCTTTGCC TGGGTTTTTG AATCAGGGCC CCAGATAGAA CAAAAATGTG GTTATTCAGT
49501  TGTTCCACCA TCACTTGTTG AGAAGACTCT TTTTTCATTG AAGTGTTTTG GCACCCTTAT
49561  CAAAAATCAA TCTACCATAA ATGTGAGAGT TTATTTCTGG AGTCTCAATT TTATCCCATT
49621  ATGCTATAAT CTATAATCCT ATCTTTTTTT TTTTTGACA GAGCCTCACT CTATTGCCCA
49681  GGTTGGAGTG CAGTGGCCCA ATCCCGGCCA CTGGCTCCTC CTCCCAGGTT CAAGCAATTC
49741  TCCTGCCTCA GCCTCCCAAG CAGCTGGGAT TACAGGTACC TGCCACCATG CCTGGTTAAT
49801  TTTTGTATTT TTAGTAGAGA CGGGGTTTCA CCATGTTGGT CAGGCTGGTC TGGAACTCCT
49861  GACCTCAGGT GATCTGCCCA CCTCAGCCTC CCAAAGTGCT GGGATTACAG GCATGAGCCA
49921  CCACACCCAG ACTATAATCC TATCTTTATG TCAGGACTAC ACTGTCTTGA TTACTATAGC
49981  TTTTTAGTAA ATTGAATTCA AGAAGTTTCT CAACTTCAAA TTTGATCTTT TTTTGGAAGA
50041  CTATATTAGC TATTCTCAGT CTGCTGAATT TCCCTAGGAA TTTTAGGATC TATTATCAAT
50101  GTCTATTCTA TTTTTGTATA TGTTTTAATA TTTTCATAAG AAACTTTTTT CATTTAAACT
50161  TTTTTTTTTA AGAAAAATAG TGAAAATCAG AATACTGGGG GTCAGGCGCA TTTAACAGGC
50221  AGAAGAAGAA TAAAAACTTG TCATATAAAC AAAAAAGAAA TGACCAATCA CATTGTGGAA
50281  GCCATGGAGT GGTTATAGGT GCCAAAGGCT GCAGAGAAAT GGTGTCAGAT ATACCTGAAA
50341  ATTGTCCATT GTATTTGGCC ATTAAGAGAC TTAGAAGACT TAAGCCATAG ATTGCTCAGT
50401  GAGACCCCGA GGGCAAATGG TCTGAAGGTG AATAGATCAT TTCACCTTTA AGAGAGCAGG
50461  TAGGAAGCTA TAAATCCAAG ATTAAAAAGT TGACTGAACT GTTAAAGAAG AAACTCTAAT
50521  CTTGAGCCAC CCTATCCTTG CTCCACCTTC TGCTGCAAGC AAACAGAAAT GCTGAAATTC
50581  AACACTCACA AAGGCTGGTA AGCTGGAAAT GACAAAAATT ACTCCTGGGA AAGTCAGATT
50641  TAGAATTAGG CCATATTTGT TGGGGTTCAG ATTTTCATGT ACACTTGGGA AAGGGTTTAG
50701  CTTATAGGCA CATGCATGAA GGGAACTGGT ATAGGGCTGT GTTCATAAGG TCAAGAGTTG
50761  AAGGCCAGGC ATGGAGGCTC TTGCCTGTAA TCCCAGCACT TTGGGAGGCC GAGGCAGGAG
50821  GATGGCTTGA GCCCAGGAAT TCAAGACCAG CCTGGGAAAC ATAGGGAGAT GCTGTCTTCA
50881  CAAAACAATT AAAAAATAAA ATTAGTCAGG TGTGGTGGCA CACACTTGTG GTCCCAGCCA
50941  CTCAGGAGGT TGGGAAGATC ACTTAAGCCT GGGACATTGA GGCTGTAGTC AGCCATGATA
51001  GTGCTACTGC ACACCAGTCT AGGTGACAGA ATGAGACCCT GTCTCCAAAA AAAGAGCTGT
51061  ATCCACATCC CAGGAAAGTG GTTGAAGATC TACTTTTCTC TGTAAACCTA ATAAAGAATA
51121  GAGTGACAAA TGTGTGTTGT GGAAAGAAAT GGGGTGAGAG CTACGTAGAT GCAAAACAAT
51181  ACATCCCCAC ATACCACTTG TTAATCATCC TTTTCCACCC ACTTATGGGA TGAATTGCAT
51241  CTCCCCAAAA GATACTCTGT CCTAACCCTC AGTACCTGTG AACCTGACCT TATCTGGAAT
51301  ACGGTGAGTT CACTGGTTAA GAAGAGATTA TAGTGGAATA GGGTGAGTCC TCCAACCAAT
51361  GACTGGGGTC CTCACAGACA CAGAGGGATG ATGGCCAGGT AGAGATGGAG CAGAGATTG
51421  GAGTTATGCT GCCACAAACC AAACACAGGA AGCTGCTAGA AGTGGAAACA GGCAAGAAAG
51481  AATCCTTCCC CAGAGGCTAC AGAGGGATCT TGGCCCTGAT AATACCTTGA TCTCAACTGG
51541  CCTACGTAAC TGTGAGAGAA TAAATTTCTT TTGTTCTAAG CCACCCAGTT GATAGTACTT
51601  TGTTACGGCA GCCCTAAGGA ACTTGATATA CATTTCTTTT ACTGTCATAG AAGTTTTGAA
51661  TCTTTTAAGT AGGTCTGTAC CCTTCCTCCC AGTGTCAACG CATGGAATTC CTCTCCTTGT
51721  GCCTTGAAAA GTGAAAGGTG TTTGAACTGG TAATGAAAGA AATCTCAGCA TGAGGCCAGA
51781  TGCTGTACCT CACACCTGTA ATCTCAGCAC TTCGGGAGGA TGAGGCGGGC AGATCACTTG
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51841  AGGTCAGGAG TTCTAGACTA CTCTGGCCAA CATGGTGAAA CCCCATCTCT ACTAAAAACA
51901  AAAAATGTTA TCCTAGCCGG GCATGGTGCC TGTAGTCCCA GCTACTCAGG AGGCTGAGGC
51961  AGGAGAATTG CTTGAACCCG GGAGGTGGAG GTTGCAGTGA ACTGAGATCA CGCCACTGCA
52021  CTCTAGCCTT GGTGAGAGAG CAAGACTTGG TCTTAAAAAA GAGAAAAGAA AAATGAAATT
52081  TCAGCATTAT AGAATAAAAA TGTTTCCCCT TCCCCCCAAA CTTTAAAAAA GCAGAAGTCT
52141  GCATCATAAA ATGGTCTTTG CCAATGTTAT TTTTATTATA ACAAAGGAAT CTTGCAAGGC
52201  TACCAGATCT CAGCAATTGT CACTATGTTC TGTAAAAATC ACTTCCTAAA ATGTCTGAAT
52261  TGACTGCTTG TCTCATTTAT TTGTTTCTCG TGTCATACTG CAATGGATAT CTGTCTTGTT
52321  AGTATAAATA TTTGTGCATT TTGTTGTTGT TAAAACAGCT TTTTTGGCCT GTCTTCTTCC
52381  ACCTATGAGG TAATATAAAA CTCATGTTTA ACACTTATTT TTGTAGGAGG ACAAGCTACA
52441  GACAAAACCC CTCAGACACT GAGTTAAAGA AGGAAGGGCT TTATTCAGCT GGGAGCTTTG
52501  GCAAGACTCA CATCTCCAAA AACCGAGCTC CCTGAGTGAG CAATTCCTGT CCCTTTTAAG
52561  GGCTTGCAAC TCTAAGGGGG TCTGTGTGAG AGGGTCATGA TCGACTGAGC AAGTGGGGGT
52621  ATGTGACTGG CAGCTGCATG CACCAGTAAT CAGAACAGAA CAGGGATTTT CACAGTGTTT
52681  TTCCATACAA TGTCTGGAAT CTATAGATAA CATAACCGGT TAGGTCGGGG GTCAATCTTT
52741  AACCAGACCC AGGGTGCAAC ACCAGGCTGT CTGCCTGTGG ATTTCATTTC TGCCTTTTAG
52801  CTTTTACTTT TTCTTTCTTT GGAGGCAAAA ATTGGGCATA AGACAATATG AGGGGTGGTC
52861  GCCTCACTTA TTCACCCCCT TTGAGAATCT CACTCATTAG TGGGAGTTCT CACTTTTATT
52921  CTCACTACCT ATGTCTTCTT GAAAGACAGA TTGATAATGA TTCATATAGT ACACTTGTGC
52981  TGAAGCATTT TGGTGAGCTA AGGTAGTGAT GAAGCTTTTT ATCATTGGA GAAGTACAGG
53041  TAGCAAACAA GGAAGCAGTA AGCAGGTTTC TATTAATATT ATAACTCCTA TTATAAGAGT
53101  TTTAAATCTT CTTAGCACTC GGAACCATTT TTCAAACATG GCCCCAGAAA CAAATCCATA
53161  CCACACCTAC ATGGGCACAT GTGCCACTTT TGTCATATTT CTAACTATGT CTTCAACTAC
53221  TTGCCCTTAA TCATCTATGT GTAGACAGCA ATTAGTAAGG TTAAATTTCC TACAGACCCC
53281  TCCTTCAGTT GCTAGCAAGT AGTCGAGAGC CAATCCATTT TGATAGATAG CATTTTGCAT
53341  CTGAGTTTCT TGCCAGGCCA CAGTAGTCAG GGCTCTGCTG GTCTTATTAG TAATTATTTC
53401  TAAGACAGCT TGTAACCGTA TGATTCAGTT GAGCATGTAA ATGGGGGTCC CATATCCCCA
53461  CAAGCCGTCT TGTGCCCAAG TAGCAGGCCC ATAATATTGT ATGATTCTCT CAGGGGGCCA
53521  TTCATTATTT TTCCAATTTT CTATAGCTAT GCTTTTTTTT TTTTTTTTTT TTTTTTTTTT
53581  TTGCGGGAAG CATATACAGG GAAGCCCAGG AGTTTGCCTG TCTTTATGGG CAGTAGGAAG
53641  AAAGATGGTT TAATAGTGTC AATAACACAA CTACCTGCCC ACTGGTCAGG TAATTTGGCA
53701  TAAGCTGTAT GCCCACATAT CCAGTATAAT CCAGTGGGGG CTGTCCAGTC CCGGTGGGAC
53761  TCTGGGTGGG TCCACACAGT TTGCAACTTT GGGAATTTAC TAAATAGATT TTTCTTAGTG
53821  TGGTTTGAAC TCCACTAGGT GGCTGTTTTT ATAGTACTAT TATACAGTTT TTGCCCAAGG
53881  CAGCTGAGTC TTCCCACAGG AAGGGTGAAG TCCTTCCCCA CTTTTGCTAT ACAGTATTGT
53941  CTAATGATTG AGGCTTTTAG GACCCAGAAG TTATCAGGGT GAGTCTTTTG AGCTGGGAAT
54001  TTATCAGGAA CTGGGTCTGT AGGTACTAAT TCTCGTGCTT CCCATGGCCA TTGATCTCCC
54061  ATTACAGTTC CTCCACATAC ATACATAACA TGAAGTGACA TTGAGAGACT GGGCTACATG
54121  CTCAGCTAAT TGCAAAAACA AATTTCTTGT TTTTCCTGGA ATTTCTAGTA CTGGCACATT
54181  CAGTTCATCA TAAGAAGGTT TGAAATACTG GCTCAGGGA GCATTTATAA ACTTCTCCTC
54241  AAACCACCAT ATTTACTCAA GGATCCAGTC CAGCCCCAAC TATTTCTAAG GTTACACGAT
54301  CCCCTTTTTT CCAGTGAGAA TCAAGGGGGT TGGTTATTAC TAGTTCTAAG GGGTTACACT
54361  GACCACTGGT ACAGGAAGGG CCACTTTTCC CTTTCTGAAG GTGGACAGGA TTCTTTTTAT
54421  TTTTTAACCA AGTTGCCTAA ATGACACAAG ACCAGTATCT ACATTTATTT CCACGCAGTC
54481  TTAATTCATG ACAAGCGTAC TTATTTTCTG CCATATAGCC TCTTTCCTAA TGAACAGAAC
54541  CACATCCTAT TTCTAACTTA TTACTATTAA TGACAGCACA GGCATCAAAT TTCAAGGTGA
54601  CTTGTTTGGG CATTCCTTTT TCTTCTGTTT TGGCTAACAC TTTACTCGTA TCGTTATGA
54661  ACCCCCACCA GTCCTCAGTC CTCAATCTTA TTTCAAAAAC TGTGGTCGTG GGAGGCTCAG
54721  ATGGGTCATA ACACACATCA GGTTGGTCAT TTCTTGGGCT ACCTGCCTTG TATAGAATAG
54781  CATTATACAA ACAAGTTATT TTTAGAGTCT TTGTACACTT ATAATAACCA TAAAATAATA
54841  AGACTGTAGC AACTTTTTGT CCTACCTCAG TGACTTGATG TATACACTGG GAACAGCCCT
54901  CAGTCTGAGG AAGGTTAGTT GAAGTCTTTA CTGTGCAAGT CCAAATTTTA AGGAAAATGA
54961  GTCCCTTGAT GAGTTTCTC ATGTTTCGGC CATGCATGGA CCAGTCAGCT TCCGGGTGTG
55021  ACTGGAGCAG GGCTTGTTGT CTTCTTCAGT CACTTTGCAG GCGTTGGCGA AGCTGCCACG
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55081    TACAGCTCAC AGTCTACTGA TGTTCAAGGA TGGTCTTGGA AGTTGGGCCC ACTAGAATTA
55141    ACTGAGTCCA ATACCTCTAC TCAGTCACTT TCAACTGGGC TTTCTGATAC CAGGAGCAAG
55201    GTGGCAGGTT TTAGGGTGTT GCAAATTTCA ATGGTTATGC AGGGATTTTC ACATAGCAAA
55261    CTTTGGTACT TGGTTAATCT AGCATTGTT AGCCAATGAT GTATTTATTA AAGTCACCAC
55321    AGCATGGAGG GCCTTTAAGT TTAGGTTTTG TCCAAGAGTT AGCTTATCTG CCTCTTGTGC
55381    TAGCAGGGCT GTTGCTGCCA AGGCTCTTAA GCATGGAGGC CAACCCTTAG AAACTCCATC
55441    TAGTTGTTTG GAGGCCCAGC CTCGGCCAGG GCCCCACAGT CTGGGTCAAA ACTCCAACCG
55501    CCATTTTTTC TCTTTCTGAC ACATAGAGTG TAAAGGGTTT TGTCAGGTCA GGTAGCCCCA
55561    GGGCTGGGGC CGACATGAGT TTTTCTTTTA ACTCATGAAA AACTCATTGC TGTTGGTTGT
55621    AATAGATGTA GTTTATCCAA TCTACATTTT TATTAACTGT CACCCACCAA AATATTGACT
55681    CAAATCCTGC AGCTATTTGA TTTTGGGATT TAAATTGATC TGCTATTCCC TGTGGGACTC
55741    CAATTGCATC TAAATAGATG TGAGAGTTGA AAGACACATA AGGGTCTTCT CTTGCTTTAC
55801    GATGTCTTAT TTTTCCTCCC TCTGGTTGAT GAAATGCTAG GGTGAAAGGG ATAGCCAACT
55861    GGACTAAAGT ACAAGTGCCG CTCCAGTTAT TTGGCAGAGT GCCCAGTAAA GGTCCACCAC
55921    AATACCACCA CACATCCGCT TGGGGATGAA CAAAGGCTGA CTGATTGAGA AGCTCCTGAA
55981    AATTCTTAAG CTCACTGCAT CCCTTCAGGT CTCCAAGGAA TGCTAAGTTT CCTCCCTGTC
56041    ATGAGAGACA AGAAGTGAAC TTAGTTTTGG GAGATGGAAG CTGGATGGCC CTCAGGGGTT
56101    GACCTGCAGG GTGCTGGACT TTGGGATATA GCAGAGAGAG CTTGGCACGA CTTATTACTC
56161    CAGGCTGTAG CATCCTGGAA AACAGTTACC ATGCAGCCCA TGCCTGGTCA ACAGGAGGAC
56221    CACCTTAGTG GAAAGGGGAT AATCTGGCCC TCTGGCCTGC CATGTGCACA AGCATAACAA
56281    TTGGTTTTGT TTAATGTGTG GACAGAATAT TTGATCCATT CCAACTGGGC ATTTGCATCT
56341    TGGTATCCTG CTTAATTATC AAAGTTTGTT TTAAGTCTTT AACTTCTATG ACCCTCTAGT
56401    AAAATGAATG TATGATTTTA GGAAATTACA AAAACCGGTT GGGGCAGTCC ATCCTCGCTC
56461    TTTAGTGGTC CACACAACAT TCGACCAACT ATGGCATAAA AGCTCTACAT CAGGGGGCAA
56521    GACTCCTCGT TGACACTGGG GTCTTTATTG AAATCTCTCT GGATTAAATG GTCTCAGTTT
56581    ACTAAGGCTC AGTCTGAGGA GAGTCAGGAG GGACAGAGGT ACTTTTCTGA AGTACAGAGA
56641    TGTCTTCGAC TTGGCAAGTC CCCACAGGGT ATAACAAGGC AAGCATTAAA TTCAATAGTT
56701    TGAGGCAAAA TTGACTTGGT TATGTTAATA ACTAGATGGT CAGAAATAGA GTGAGGGAAG
56761    AAGAAAGAGT AATAGAATAG ATGAAGGAGT TAAATTTTTC TTAGCTTTAG TTTGGTAGGG
56821    TTTTCCCCTG GGACTATGGC CCATGACTCT GGAGGGGGTG GCACTTTCTT GACTCGGGTG
56881    TGATGAGTCC ATCCCTTTTT CACCGTATGA ACAACAGTCT CGGTGGTTAG CAGCACAAGG
56941    TAGGGTCCTT CCTAGGCTGG CTCAAGTTTT CCTTCTTTCC ACCCTTTGAT GAGAACATGA
57001    TCTTCAGGCT GGTGCTGGTT TACAGAAAAT TCTAGGGGTG GTACATGTGC TAAAAGACTT
57061    TTAGTTTTGA GGGAAAGGAA AGTGGAAGAT AAACCAAGTA TATAACTTTT AAGAAGTTGA
57121    CCTTTTGTTT TAAATGTGGG GACATCAGCA GTGGACTTTA TAGTCCTTGG TGCCTTCTTA
57181    CTGAGAAATT TCCTTTAGCA CCTATTTTTA TTAGTTTTTA GACCAAAGAA AGTCAAATGC
57241    CATTTTATAT TTGACAACGC TTCTTGTATG TTTATACCAG ATAAGCTAGA TTTCACCTTT
57301    ATATTGGTGT GTTATTAATG TTAAACTTAG TTTTAATAAA ACTCTGTAGA CATATTTATT
57361    TGATTTTTAA TGTCTGACCA TAAGGTAAGA TTTTTATAGA CTTTTCTTTA ACCTTTTATA
57421    ATTTTTGTTA AAGAACAGGT TAGTGCTTTA AGAAAACCC GTTGTGTTTT TATTTTAATG
57481    TTCAGTTCAC AGAAAAACTG TATGATACCC CTTAACTTTA GCCAATATGT TTAGACACAG
57541    AATTTTCTTT ACAATTAAGG TTTCAAAACT TGCTTAAACC TTCAAAACAA TTTTTGTAAC
57601    CTTTTAATGT AGGTAAAAAT CCACATTCTT ATGCATCCTC ATAATCCTTT TACCAAAGGT
57661    ATATTTACT TTCCTTACAT ACCTTGCACA TAAACTGTTT ATTCAATAGT TTTACATTTA
57721    GAAGGAGGCC TAATTACTTT TAAATTATAC AACATTTCTT GCATAAATTT ATTTTTCTAA
57781    CACACATTTT TTTCATGACT TTCACAGACA ATTCTTCGAC ATGCCTCAAC TTTCTGACTT
57841    ATTGCAAACA TCCCTTTCTT TAAACAACTA GTTAATTTAT CTCAGGACAA GGATTTTCCA
57901    TACAACATTC TTTTTTATAT AAATTCTGCC TCCTCTTTAT TTCCTTTTTT TTTTTCCGAG
57961    GATGATAACC ATTCTTTTCC AAAGCGAACT TCTTTTATGT CTGTGGACTA GACTGTCTAA
58021    GGCCACAAGA TTAGAAGTTA CTATAATACA TGTTACACTG TTAACTTTTA GCAAACTTTA
58081    CTTTTGTTGA AAACCTTGTA AGTTGGGAT TTCAATTATC CTTTGCTATT AATAAGACCT
58141    TATTTAGTCC AAATTAACTT AGAATTGGTA TAGATGGCTT TTTTTTTTTT TTTAATTACC
58201    TGGGAGGAAC CATCTATCCT CCTGTCCTGA AGGGAGTTCC TCCTAGGTCT GGTCAGAGCT
58261    TTGTATGGTA ATTAAGATTT AGATCCCCTG TTAGGAAACC TGCCGGGTTA AGAGAATTTT
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58321  CAGTGGTTAA  TGTTAAATCA  TCTTCTTTTT  TCTTTTTTCC  TTAGGATACT  TCTGAACCGG
58381  TGAGGTGTGC  TCACAATGAG  GTTTCCTGTA  AAAGTTATTT  TTTTACTTTC  TTCTGTTAGC
58441  AAAGCAGTTG  CCGCTACAGA  TTGAATGCAT  TTGGGCCATC  CGCGGGTTAC  TGGGTTAAGG
58501  ATTTTTGATA  GGAAGGCCTT  AATGCTTTTG  GAATATGCCC  TGACAACAAA  GTGCCAGTTC
58561  CTTCCCGGTG  TTCAGCCACT  GCGTTGATCC  TCCACGAGGG  CCTGCCACGT  GCTGCTCTGG
58621  TGAGGCGTTC  CACCGGGGCA  ATTGCCTACC  TGGGAGCGCT  CTCCAGATCT  GTGTCGCTCA
58681  AACTGGCTGG  AGTTCCCCGT  AGGGATGCTC  CACAGGGCAG  GCCTAAGTCG  CCTAAGGGGC
58741  TGCCTTGACC  GTCCGTTAAT  CACCTCTGTC  TCCAAAAACC  AGCTCCCTGA  GTGAGCAATT
58801  CCTGTCCCTT  TTAAGGGCTT  ACAACTCTAA  GGGGGTCTGC  ATGAGAGGGT  CGTGATTGAT
58861  TGAGCAAGCA  GGGGGTACGT  GACTGGGGCT  GCATGCATCA  GTAATCAGAA  CAGAACAGAA
58921  CAGCACAGGG  ATTTTCACAA  TGCTTTTCCA  TACAATGTCT  GGAATCTATA  GATAACATAA
58981  CCTGTTAGGT  CAAAGGTCGA  TCTTTAACCA  GACCCAGGGT  GCGGTGCCGG  GCTGTTTGCC
59041  TGTGGATTTC  ATTTCTCCCT  TTTAATTTTT  ACTTTTTCTT  TCTTTGGAGG  CAGAAATTGG
59101  GCATAAGACA  ATATGAGGGG  TGGTCTCCTC  CCTTAATTTA  AACAAAATTT  TCAAAGTCCT
59161  ACCCCAAGTA  AATTGGCAAA  TATTAATAAA  GTTATGGCAT  AGAAAATAAA  AATGATTGTA
59221  AAAGGCGTAA  AGATATTTCT  GTGGGGAAAA  CATTTGTTCA  TTAGTTATCA  GTTAAAATTC
59281  TGTGAAAAAT  AACCACTAGA  GACCCTAAAG  TACCCAGGGG  CTAATAATAA  GAAGGGAGGA
59341  ACACCCTCTC  AGTCCCCACC  GTTACCTCCC  CAGAAGGGAA  GAGGAAGAGG  GTGACTCCAG
59401  GAGAGCTGTG  GTCTCCCCTC  CCCATATGTC  CACATATACC  TGACCTCCCC  TCCCCAAAAT
59461  ATATACCCAA  TATCTCTCCC  ATATATACAT  ATTTATCTGA  CCTCTCCACA  TATGTATACC
59521  TAAACTTTCT  CTATATATCC  ACATATACCT  AACCCTCTCA  CACACATATA  GCTGACCTCC
59581  AGTGGAGGAA  AATGGGGAAG  AGAGAAGAAG  TTATCAAAGG  ATAAATCTAG  GTCATACTCA
59641  GAAATGTGAA  AAACAAAAAC  CACACACAGA  AAAAAAAAAC  ACACACAAAA  AAGAAATTGA
59701  TAAATTTGTT  TGTGTCAAAA  TTAAGAATTC  CGGTTCAATG  AAGGATCCCA  TGGATAAAGT
59761  TAAGACACTG  CTGTAAGGAT  GGTAGAGAAT  TAAATGTCTG  AATCAGACGA  AAGGATGAGT
59821  AATTAGAATG  CACAAGGCCA  AGAAGAACAA  AACAGAAACT  CCACATAAAA  AATGTATGAG
59881  GCCGGGCGCG  GTGGCTCATG  CCAGTAATCC  CAGCGCTTTG  GGAGGCCAGG  GCGGGCCGAT
59941  CAGGAGTTTG  AGACCAGGCT  GGCCAACATT  GTGAAACCCC  ATCTCTACAA  AAAATACAAA
60001  AAATTAGCCG  GGCGTGGTGG  TGGGTGCCTA  TAATCCCAGC  TACTTGGGAG  GCTGAGGCAG
60061  GAGAATCACT  TAAACTCAGG  AGGCAGAGGT  TGCAGTGAGC  TGAGATCACA  CCATTGCACT
60121  CCAGCCTGGG  TGACAGTGTG  AGACTCTGTC  TCAAAAAAAA  AAAAAAATTA  TATATATATA
60181  TATATATATA  TATATATATA  TATATATATA  TGAAATAAAT  GAACAAGAAA  TTTAGATACA
60241  GGAAAATCCA  AAGCACTTGG  TAATGAAAGA  AAGGTAAAGT  GATGTGTCCT  TTTGCATTTA
60301  AAAGAGAGCA  TTAACAAATT  AGAGAGCTGA  ATAATGCTCA  GTATTGGTGT  GGATATGGAG
60361  ACTCAGGAAT  CCTCATACAC  TGCTGATGGG  AGTGCCCACT  CCCTGGGAAT  ATTTTCCAAA
60421  TATCATCTCA  AACATATCCC  ATAAAGGTGA  CAGGAAAGTG  TGGGCTGACT  GATATCCTTC
60481  ACTGAGAGAG  GTGGAGGTAA  AATGAAGTCA  CTGCACAATA  TAGAGTTGGA  AGCAATGGAT
60541  TAGATGTCCA  CATAGTTACG  TGGAAGAATC  CGTAAGATAC  ACACACACAC  ACACACACAC
60601  ACCTTTGTGT  ATATTGTTCC  TGGCAGGTAG  GCATGGAGGT  TTAGAGGCTT  TCTACATCAC
60661  ACCTACTGCA  CACAGTAAAT  GGCCAGGCTG  AGCACTGACT  TCCATGAAGG  GAGATTGAAG
60721  GTAAGAGATT  GAAGATTGTT  CCCTGGTCTG  GGACCCTGCA  ACTGAATATG  CAGAAAAAAG
60781  TACACCCCGC  CACCCCGCTT  CCCATCTTTC  CTACCTGATT  AGAATAGCTT  TTTCAGAAAA
60841  CGTTGGCCAG  GGGTTGTGGC  TCACACCTGT  AATCCCAGCA  CTTTGGGAGG  CTGAGGCGGG
60901  CAGATCATCT  GAGGTCAGAA  GTTCCAGACC  AGCCTGGCCA  ACATGGCGAA  ACCCCATCTC
60961  TACTAAAAAT  ATAAAAAATT  AGCAGGGCAT  GGTGGCACAC  ACCTGTCATC  CCAGCTACTC
61021  GGGAGCCTGA  GGCAGGAGAC  TCACTTGAAG  CACAGTGATG  GAGGTTGAAG  TTAGCTGAGA
61081  TCTTGCCACT  GCACTCCAGC  CTGGACAACA  GAGTGACACT  TTGTCTCAAC  AACAACAACA
61141  AAACCCACCA  AAACTTTAAA  TCTACCTATG  GCCAAATGCC  TGCTAAAATG  AGCACCCAAG
61201  AAGCAGTGTT  CAGGAAAGTC  AGATGAATAC  CCTAAAATTA  GATGCAATGT  TGGCTGGTCA
61261  CAGTGGCTCA  GGCCCTGTAA  TCCCAATCCT  TCTTGGGAGG  CCGAGGCGAC  AGATCGCTTA
61321  AGCTCAGGAG  ATCGAGACCA  GTCTGGACAA  CATGGTGAGA  CCGTGTCTCT  ACAAAAACGT
61381  ACAAAAATGA  GCTGGGAGTG  GTGGCGCACA  CCTGTAGTCC  CAGCTACTCA  GGAAGCTGAG
61441  GTGGGAGGAT  CTCTTGAACC  CAGAAGGCGG  AGACTGCAGT  GAGCAGAGAT  CATGCCACTA
61501  CACCCCAGCC  TGGATGATAG  AGCCAGACCC  CCATCTCCAG  AAAAAAAAAT  AAAGAGAGAG
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61561   AGAGATGCAA TATTTAGGGT TCAACAAGAC TGAACTTCTG ACTCCTTTCC CTACCTCTCC
61621   AGCATGTTAG ATTCTGGGTC CTTCATCCTA ACCCCTGTT  CATGCCATAG CCACCCTGTG
61681   GTACCAACTT TGGAAGCCTG GATCTTCATC CCCTCATGAT AATGAGTGTC CCATTCAGGT
61741   CTCCATGCTC AGCTTGGCAA GAGTATCTGT CTTCTCCTCA TGGGACGGTC ACATTCACCC
61801   AGCACTGACA GGTTCCATTC CCACTAGGGT GGCACCCTAT ATGGTCTGAG TCCAGGCCTT
61861   CCTGGTCCCT CAGTAATCTC AGCATGGTAG CACAATCGAA AAGGGCTAGG CACGGCAGCA
61921   CCATTTCCCA CCAAGAGGTC TGATGGCTCA TCACATAGAC TGAAGGAGAT TCTGAAGAGC
61981   AGAGGTGGAA TGAAGAATGA ATCCTGGGCT CTGCTCTTCC TAGGCCTGTC TTCCTCTCTC
62041   CCGAGATGTT AGCTAACTCA TGAGAGCCAG AAACCAACTG CAGGCTGGCC TCAGGCACTT
62101   AGGTAGTGCT TCAGCCTCAG CAGTCCACAT TCTAGGAACC CTCATAATAT GGGTTGAAGT
62161   ATGCATTCCC ACAAAAATAA AGTTGTTGAA GTCCTAACCA CCAGTACTGA AATGGGAAAA
62221   GTTCCCTTGT CCCGCTCGCA TGGCATGTGA TAGGAGTGTG GCTAATTTCT TCAGTGCCTG
62281   GCTGCTCAAA CCTCTAGGGG AACAGTAAGA CGGGCAGGTT GTGGGTCTCC AACCCCATGA
62341   CCCCACCACA GTGTCTAGGG TTGAATGTTT ACAGCTCCTG AAGCCACAGT GGGTGTGTGT
62401   TACAGGGTGC TCTTTTAGTT TTGCCATTTA TAGGCAGCTG GTGTTAACCA ACTCAATTAG
62461   ACCGTCTACC TTGTCCCAAG GACAGAAGAA GGCTTTCTGT ATCCCAGGTT CTTGCCTTGG
62521   TGTACCGGAA TAAATCAGAC CACACCTGGG CTTAGAGAAA GAGTGCAAGG TTTTATTAAG
62581   TGGAGGTAGC TCTCAGCAGT TGGGCAAAGC CAAAAGTGGA TGGAGTGGGA AAGTTTTCCC
62641   TTGGAGTCAG CCACTCAGTG GCCCAGGCTC TCCTGCAACC ACCCCAGTCA AATTCCGCCT
62701   CATTTTGCCA GGCAAACGTT TGTTGTGTGC TCTTCTGCCA GTGTGCTCCC CTGGACGTCC
62761   AGCTATTCGT GTCTTGTGGC AGGCCAGGGG AGGTCTTGGG AAATGCAACA TTTGGGCAGG
62821   AAAACAAAAA TGCCTGTCCT CACCGTGGTC CCTGGGCACA GGCCTGGGGG TGGAGCCCTA
62881   GCCGGGGACC ACGCCCTTCC CTTCCCCACT TCCATATCAT TTAAAGGGAC CATGCCCTTC
62941   CCTTCCCAGC ACTTTCCCCC TCCTGTATCA GGACCTGTGA ATGTGGCCTT ATTTGGAAAT
63001   AGGGTCTTTG CACTTCATCA GTTAAGATAA GAGTGGGCTC TAACCCAACA TAAAGGGTGT
63061   CCTTATAAAA AGGAGAAATG TCATACACAG AGACTGACAC CTATAGAGAG AAAATGTGGT
63121   GAGTAGACAC AGGGAGAATC ACCATTCAAG TCAAGCAATG AGTCTGGGGA TACCAGAAGC
63181   TGGGAGAGAA ACCTGGAACA GATTATCCCT CATTGCCTTC AGAAGGAATC AAACCTGATG
63241   ATACTTTGAT TTCAGACTTC CAGCTTCCAG GACTGTGTGA CGATAAATAT CTGTTGTTAA
63301   GCCAACAAGT TTGAGGTACT TTGTTACTGC AGCCCCAGAA AACTAATACA GTAGGTACTA
63361   TGGACTGAAT TGTGACTCCC CGTCGCAAAA TTCATATGTT GAAACCCTAA CCCCCAGTGT
63421   GATGGTACTT GGAGCTGGGG CGTTGGGAA  GTCATTATAT TTAGACAAAC TCATCAGGAT
63481   GTGTCTCTCA TGATGAAATT CATGCCCTTA TTAAAAGAGA CAACAGGCCA GGTGCAGTGG
63541   CTCATGCCTG TAATCCCAGC ACTTTGGGAG GCTGAGGTGG ATGGATCACC TGAGGTTGGG
63601   AGTTTGAGAC CAGCCTGGCC AACATGGTAA AACCCCATGT CTACTAAAAA TACAAAAATT
63661   GGCCAGGTGT GGTGGTGCAC GCTTGTACTC CCAGCTACTT GGGAGGCTGA GGCAGGAGAA
63721   TCCCTTGAAC CCAGGAGGTG GAAGTTGCAG TGAGATCACA CCACTGTACT CTAGCCTGGG
63781   TGATAGAGAC TCCATCTCAA AAAAAAAAAA AAAAAAGAC  AATAGAGCCA GGTGCTGCAG
63841   CTGATGCCTG TAATTCCAAC ACTATGAGAG GCTGAAGCAG GAGGCTCGCT TTAGCCCAGG
63901   AGTTCAAGAC CAGCTTGGAC AAAATAGTGA GACCCCCAAC TTCTAAAAAT TTAAAAAATG
63961   AACTGGGTGT GGTGGTACAC ATCTGAGGCT CCAGCTACTC TGGAGGCTGA GGTGGGAGGA
64021   TTGCTTGAGC CCAGGAGGAG GCTGCAGTGA GCCATTGCTG TCCAGCCTGG GCTACACGAG
64081   AACCTGTCTC GGGAAAAGGA GAAACAGTG  AGACCTCTTT TTCTCTCCTC CTTCTCTCCA
64141   CTGCCTAAGC CCTACAAGCA CAAAAAGGAC ACCACATGAG CACATAGTGA GAATGCTGCT
64201   GCCACCAACA AGTCAGGAAG AGAGCGTTCA CCTAGAAACT GAATTGGCCA GCACCTGGAT
64261   CTTGGACTTC TGAGCTTCCA GAACTGTGAG AAAGTTATTT TTTTTTTAGC GACTAAGTCT
64321   ATAGTATTTT ATTACAGCAG CTCAAGGTAA CTAACATAGT AGAAGGGATG AATTATGGAG
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64441   ATGATGAATT ATTTTTAAGA ACTTTTAAGG GATCTGACAA GTTTGCAAGA GCTAGAGAAT
64501   GCTTTACAAC GTGATAATAG AATGCTCTGT GATGACAGAA ATCTTTCCAC ACTGTTCAAA
64561   ACTAGCTACT GGCCACTTGT GACTATTGTG CACTTGAAAT GTGACTGGTG TCTGAGGAGC
64621   AGAATGTTTA ATTTTACTTA ATTTTAATTC ATTACAATAG CTACATGTAG CTAGGGGCTA
64681   CTGGATTGAA CAGCACAGCT CGAGTCTTTT AGAGGGAGAC AGGACTCACC AAGGTGGATG
64741   CTGGTGGCCA AGCAGCAATG GCAGGTAGTA CACACACAAG AGGCAGATGA TACAACACAT
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64801   CCTTCCCAAA CCTGGAGATA AGCTCACCCC ACAATCCCGC CGCTGAAATA GAGTTGATGT
64861   TACCAATGTG CATTTTTATG TCCTTTTCCA TACAGAAAGA TCATTCAACA AGTACTATGG
64921   TACTTAAAAA ACAACATTCA ATTCATTATT ATGACAAAAT TAAATTAATA GCTCTTCCTT
64981   AAACTTTTAA ATTCAATTTA CAATGCTTAC TATTGGCATT TATTAATCTA CCAATTTTTT
65041   CCCATAGAAC CCATAGAAC  AATAATCTAC CAAATTTTTA ACATTCATTT TTGGCAAGGC
65101   TTTTGCAATT TGACGAACTT TAAGAAGAAA ACTTATAAAT TGCAATTTTT AAATCTGACA
65161   TACTGGACTT TTAAAGTATC CAATTGACTA ATGAACAAAA CTGCTCCAAA TTTTTCAATT
65221   CTTAAAAATC TTAAGACAAT ACTTAATATG GCAAATCTTA ACTTCTTAAA CTTTGTAAGA
65281   ATGCTAATCA ACTTAGATTG GTATAAAGTT GAGTTAAAAA TCACAGGATA CATCATCTCA
65341   GCTATAAGTT TTCATGAGTT GAGTTTTTAC AATCACTTGA AATGCTTAGA ATAGGAAATA
65401   CGTATAAATT ATTTAACATA AAATATTGTT ACAAAACCTC TGGAGTGTCA GTTTCTCTGG
65461   CCAGACTTTA TGCTGCAGCA CCTTTGCCTG AGTTCTTGTC CTGCATCCAG GAAGAATTAG
65521   GTACAGAGGC AAGAGTCAAG AAGATTAGTT TTCCAATAGT TCAGCTCACC TAGTTAACTC
65581   CTGTTCACAA TCTTCAAAGT TATCAGAAAC CTGCAATTGA GGGTTATAAT CCATTCTTTG
65641   CAGAGTTTCA AAACAAGACA ACATTTGTCT ATGAATGTTA AAATGTCCTA GGGTAGTCAC
65701   AGTCAAAAAC ACAATTGACA AAGAAATTTA GTCACCTCTG TGATTTACAA TAGCCTAACA
65761   CAATAACTCT AATTATAACT GATGACACAA ACTCAGATAT CAGAACTCTA GAAATCCCCT
65821   ATAATTTTGG AACACATATT CACAGTTTTC ACTGAAATAT GACCTGAAGA TCAAATATCA
65881   CCTTATTTCA ACAATCCTAT ATAACTAAAC GTGTCAAATG ATCCTGTTTA CCTCTCCTTT
65941   GGATACTCCA GGGGCCCTCT GTAGCATCCA AAAGTTAGGG GTTAGCAAAG ACAATTTTGA
66001   AGCTGTAAAG GCTCAAAACA CTTAATGAAC CTCTAGTCAT ATCTGTTCTC TACTCACTAA
66061   ATGCTAGTAG CACCTCTCAG TTGTGGCTAA GCTGGGAGGA TCTCTTGAGC CTAGAAGTTT
66121   GGGGACGCAG TGAGCTATGA TTATGCCACT GCACTCCAGC CTGGGCAACA ATGCAAAATC
66181   CTGTCTCAAA AACAAAAACA AAAAACAAAT TGCCTATGCT GTGGTTATCT CACAATTAAT
66241   AAAAAGGAAA AAAAAGTAT  GCAGTCTTTG TAGGTCCTTG GGGTTTGTTG GAACTCAGAA
66301   AACAATACCC CAAAATAAAG ACCGCAGAAG CCAAAGTTTT TCTCTGATCT TCTCCTGCCC
66361   TCCTGTCTCT GAGTCCCATT CTCCCCGGAG TCTAGCCATA GAAATGAGAA TTCCTCTTCC
66421   TCAAGTTAGG TCATAGAAAT CAAAACACCT TTTCCCCAGA GCCCAGCCAT AAAACCTAAA
66481   AATATTACTC TAACTTTCCC TCTGTTTTTC TGTGTAAAAA CTGGCCATAA AGAAATTATC
66541   TGAACTACCT TATTTGATCA TAGATCACCA GACCGCATTC CAGAGAGGAT CCAGAAGGAA
66601   GGAATGCTGC ACAGAGAGGC CAAGAAGAAT CTAGACAGAC AGGCCTTGCT GGGTTTCCCT
66661   ACTCTGTTTA TTAGCAATCC TATTTCTACA CGGCGGCCCA TACTTTGTTG AATCTAAAAA
66721   ATAAAAATGG ACAATTTCCC CTGTACATGT TAATACACAT TAATAAATTG GATATAAATT
66781   GGATAATTTA TTAATATACA CATTAATAAA TTGGATGCAG CCGGGTGCAA TGGCTCACGC
66841   CTGTAATCCC AGCACTTTGG GAGCTGAGGC GGGCAGACCA CGAGGTCAAG ACCACCCTAG
66901   CCGAAATGGT GAAACCCCGT CTCTATTAAA AATACAAAAG TTAGCTGGGC GTGGTGGCAC
66961   ATGCCTGTAG TCCCAGCTAC TGGGGAGGCT GAGGCAGGAG AATTGCTTGA ACTCGGGAGG
67021   CGGAGGTTGC AGTGAGCCGA GATTGCGCCA CTGCACTCCA GCCTGGTGAC AGAGTGAGAC
67081   TCCGTCTAAA AATAATAATA ATAATAATAA TAATAATAAT AATAATAATA ATAAATTGGA
67141   TGCATTTTAT CCTATTAATC TTCCTCTTGT CGGTGGTTTT CAGCGACTCT TCAGAGGCCA
67201   AAGAGTAAGT TTTCCCTTAG CCCCTACAGG TTCTTATGTT TAATTTGTTA CTCTCATTTA
67261   AGACATAATT AAAGTGGCTT CTCCATGAAG ATTATTTCTG CATCCATTAT TTGGTAAGAT
67321   TGGCCGTTTT CTCCTTTGAT CTCTACTTCA CACTGACCCA CATAAAACAT CACTGCCTGT
67381   TTTTTTGTTG TTGTTGTTTG GAGACGGAGT CTTGCTCTGT TGCCCAGGCT GGAGTGCAGT
67441   GGTGTGATCT CCGCTCACTG CAAGCTCCGC CTCCCGGATT CACGCCATTC TCCTGCCTCA
67501   GCCTCCTGAG CAGCTGGGAC TACAGGCACC CACCACCAAG CCCGGCTAAT TTTTGTATTT
67561   TTAGTAGATA CGGGGTTTCA CTTTGTTAAC CAGGATGGTC TCGATCTCCT GACCTCGTGA
67621   TCGGCCCGCC TCAGCCTCCC AAAGTGCTGG GATTACAGGA GTGAGCCACT GCGCCCGGCC
67681   CCGTTTTTTT TTTTTGGTT  TTTGCATGTC TTCTCCCTTT TACTGTAAAC TATTTCCACT
67741   ACCAGCGTAG TTATCATTTC TACTGCTTAA TAATTGTTTT GGGGAAGTGA ATGCATCAAC
67801   CCACATGAAT TTCTTGTCTA TTTGACAATT TATTCTCTTT AGGAATAGTA TTAACTCCTA
67861   AGGTCCTGGG AGCCAGTCTC TGTACTTGGC TGCTCCAGGG TCCTACTTCA GTTTCCCAGC
67921   TTCTCAGTAC TGTCACTGTC AATTGTGGGT AATAATTATT TTTGTCCACC AAAAGACTCT
67981   GTATGTGAAT GAGTTTTGAA ATCTGCTGAG TAATACAGTG TCAACCCAGT TAATGATTTG
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68041   CCGGGCGGCT TGATCAGGGG CTGTCCAACT ACCGGCATTT TGATTTGGAG CGTCATCTAG
68101   TGTCTGAAAG CACAAACAAC ATCCTACATT GTAAATGCCT TTGGCTACAG AGATTGAAAC
68161   CAAAGCAAAC CTATGTTTTG AATTGTTATT CTTCAGCAGT TCTGCTAGCC TTGAAAAATC
68221   TAAAAGTTAA AAAAAAGCTT TATATTTCAT TTTCTGCCTA AACTCTTTAA AATTGCTAGT
68281   TGACAATTAG ATATTTTCAA TTTAATGAAA TTTTTTTTTA GTTCACAGAT TAATACACAA
68341   TGGGGGAGGG TTCTTATTCT GTTGGACTTT TACATAACCT CCACTTTAGT GCAGTCTGCT
68401   TTATGGGGTC TTGTTTGAGG TGTGTGTGTG TTTAAGGGAA TGTGGTTTAC AATCAAAATA
68461   TTGGGTTGCT CTTAGGCACA TTGTAAAGTC ACACACCTGT ATTCTTATTG ATACATAATG
68521   ATTAATAACA TTATTATTAC AGCCTGATCA CCATCATTAT TGATATATCT AAATAATGAA
68581   TTTTATAATT TTGCTTCCTG TCAGGCAAGA GCCAATTTCA GTGCTACCAT GTTTGTATAG
68641   CAGTATTTAT GTCTGTCATC CTCAGTCATT TTACTTCACT TGTTCTTAGC CAAACGGCCG
68701   AGAAGCGATG GTCATTTTAC TTCAAAAATG AAAAGAATTA ATATTTTTAC GTTTCCCTTA
68761   AAGACCCTAT GTTTAACCTC CACTCCCGGG TAAAATGGTC TAGTCCCTCC TTTTCATATC
68821   ATCTCTGATA TCTTTTGCAC AGCCACTATT ACCTACCGTT TTCTAGATCC CTATTCTTCA
68881   AACACCACCA TGAAGGTAGA GCCTGTCTGA ATTATTTTCT TGTCCCGTGA ACTCAGTACA
68941   TTGTTAGGCT TCTTGAAGAT GTTGATCAGT TGTTTGTGGA GTGAATGAAT CAGCTAGCAT
69001   GATTTTTCTA GACCACTGAG ACAAGTGTCT AAGACACTTG TTCCTTCCCA TGTTCTTGCC
69061   TGCCTGTGCA ATCCATGCAG TCTCATGGCT TCCCAGTGCC TCAGAATTAT CCCCTGTCAA
69121   ACAGGCATTA TAATTTCTGT CCACTGAAAA GGACAAAAAA CTAAGTGTAT AGCTAGAAGT
69181   TAAAAATTAC CGGCCAGGTA CTGTGGCTCA CTCCTGTTAT TCCAACATTT TGGGAGGCTG
69241   AGGCGGGCAG ATCACCTGAG GTCAGGAATT CGATACCAGG CTGGCTAACA TGGCGACCCC
69301   GTCTCTATCA AAAATGTAAA AGTTAGCCAG GTGTGGTGGC TCGCACCTGT GGCCCCAGCT
69361   ACTCAGGAGG CTGAGGCAGG AGGATCGTTT GAGCCCTGGA GGTTGAGGCT GCAGAAAAAT
69421   AGGAATATAC TCTCTTTCAA GAGTTCGTGG TTTTGACTGC CACCTAGCGT ACATCAGAAA
69481   AACCGCATGA CATAGGAAAT GCCTGTGACA GAGGGGTAAG GTGAGAGAGG TTGATGAAGA
69541   ATGTATTGAA GGAGTGAAAA CGCTTCCATC CCTCTACTTA CTAAATATAT TAGTTAAGTA
69601   GTTGGGGCAT ATTTTAATTC ATGCATTTTG TAGATAGAAA AACAAAAGTT TTATTCTGTT
69661   TGATTTAGTT GATACTTTAA TATGTGTGTG TTTAGGATGC ATGATTTATA ATCAGTCTGC
69721   AGCACTTCTT GGAGAAGTCT GAATTCTCAT TCTCCATTTC CTTATTGGCA ACGTGAGAAT
69781   GATTACAATG GTGGTTGTCT CATAGAATGC AGGGAGTCAG AATGAAAATA GTCCATATAA
69841   TGCCTGGTGC AGAGGAAGGG TTCAGTTAAC TGTCTGTATT AATATTACTG ATAACAGTCA
69901   TGACAAACAA AAGCTTAACA ACAACACCAC CAACAACAGT TGCAGAATTG AGCCACCAAT
69961   TTGCACACAA GATTGTAGGT AGGATGTTTT AGAAAGTTA TTATTTAATA TATGTATATA
70021   TTTTTGTACT TAAAATATGT CAGAGGTTGT TCTAAGAACT ATTTAAATGT TAACTCCTTA
70081   ATCCTCATAA TGACCCATGA AACAGGTAGG CTTATTATTG TCTCTTTACA TGTGAGAACA
70141   CTGAGACACG AAAAGGTTTA TTAACTCACC CAAAGTCACA CAGCTGGTAA AACGGCAAAA
70201   TTGAATTTGA ACTCAGACAT TCCAGGTTCC AAGACAGTCT AATTATTCTT TTGACTAATA
70261   TACTAAGCTG CCTCTGTATT TTTCCTTGAT TACTTTGTAA · AAGTATGAGG AAAATATAAG
70321   TGCTTCAAGT AACCATGAAA AATATAAACA ATCTATGTAT CAACTGAAGC ATAATTACAA
70381   ATCCTTTGAT AAGCAAACAT AATAAAAATT TGATATCAAT CAAAACTTTC ATGTAATGTA
70441   AGCAGGTTGA GATGAATTCT ATAGTAAAAA AGTGCAGAGT GCTGGAATAC CATGCTCCTA
70501   ATATATTGGC TAGGCACACC TGCCTGCTAT CAAAGGTATG CACACACCTT GGATACAGAA
70561   AGTTGGGACT GGGTAGTTAT GTGAGTGTCA TCAGAATTCT TTCCCACTTG GGAAAGAATT
70621   GTCCATCATA AGCTTGGATG ATGGACAAGG AGTGAGCTCC CAGAACAGTG ATGTGGGGAT
70681   ACATCCTCAC ATCACAGTGA GAATGAGTGT TCTAGACTGT TTACACACCT ACCACTCCTA
70741   AATGCACACA TATAATTGCT TGCACACACA CACATACACA CTCATCTCTT CTCTGGTGGT
70801   CCAGCTCTAT CTCTTATCAT TAGGCTTCTT GGGGCTAGTA CCTAGGGCCT GTATCCTTTC
70861   AGAGGCAGCT AAGGGAAGCA CACATAATTA GAAAGAATGA ACCAGCTTGT TGGATTTGGT
70921   CTCTTCGCAT CCAGCCCTCC AAGTTAAGGA GAGTACCATC TTTCTTAGGG TCACCAAAGG
70981   AAAAAAAAAA AAAGAAAGA AACAGAAGGA TATCATACAG CAAGGATCTA ATGCAAATAT
71041   GCCTCAAATG AGAGGCTACT GTGTGCTGAT CCCAATCCCA GGAACTGTAT GCACATTATC
71101   TAATTTAATC CTCACTGTAT TTCTGGGAGT ATTATTCCCA TTTTACAGAG AAGGAACTTG
71161   GCAGGGTAAC CAAGCTCATG AATGGAGAAA CTGGGATTAA ATATAAAGCT TCCTTGCTCC
71221   AGAACTGCTG TCTTTCTGCT CTTCCACACT ACCAGCTCAG CTGTGCTCTC TACATGCAGG
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71281   CAGTTTTACA AGTTTCAGAT TAGCCTGGGA CTTCCAGGGT TTTGAATGGG TTAGGGAATG
71341   GGGAACTTTT GGGTTTACTT TCCATTTTTT CTTCATACAT ATGTAATATA TAACATAAAT
71401   CTATGGTATA TATGATAAAT ATATGGCTAC ATATGAACTA TATAATCACA TATATGCATT
71461   ATAAATAAAT ATTAATTTTA TAATATTTTA AAGGTTATCA AATAAATATT AATATAAATA
71521   ATTAAATAAT TAATACTCAG CTTTGTTTTC CAAAGTGATA AATGCCTATA TTTAGCAAAA
71581   TATTTTTTGG AGGCCTGATA GTTTTTAGGA GTGTAAAGAA GTCCTGATAT CTAAATGTTT
71641   AAGAACCACT ATTTTAGGCT GTTGTCTTCT GTCTTATTTT CCCAGCTAGA CTGGTAAATA
71701   CTTGAAGGCA AACGTTTAGC CAGCACATTA ACATTTTATG TTTTTATTCT TTTGTGCTCT
71761   CAGTGGCTGT GTCTTTTCTA TCGATTTCTC ACACTGTATG ATGGTTATAT TTGTCTGTAT
71821   CTGTCCCACC AGGTATAAGT TCTTGAGAGG ACACACTGCT AGGCTGATCT TAGTTTTTAT
71881   TATTTCTCCT GGTGTCCTGT GCTTAACAAG TGCTCATTAA GTGTGTAAAA ACACAGCACA
71941   GTAAAAAACT AGACATTAAA AAATAATGTC AACCAATCTA TTGAAATTTG CATTTCCATG
72001   TTTCTTCCAA TATAGTCATT GTGTCAGGTT ATGTACTTAT TCTGATGAAG ACTATTGCCT
72061   AATATACGTT TGCATCTTGT GCTTTATAAC TGCCTTCATA TAGACACAGA TTGAGAAGGT
72121   GTAAAAATGT GCATATCCTC ACAATTGACA AATTCTTATC CTTTGAGGGT AGGTTTGACT
72181   TTCTGAAATG CTTTGACATC ATTTGAAAGA AGCTTGAAGA ATAAGATAGC TGTTAATGAC
72241   CCAGTTTCCT ATGTCACTTA TACAATTATA ATGGCAATTT CAAAATGTTA GGTAAATATA
72301   TTTTGCAATA TATTGTTCCT TTTGTAATAC TCTCTATGTA TTTATTTATA TTTTTAAATT
72361   TTATATTTAT GTATTTATTT TTCTGGACAG AGTCTTGCTC TGTTGCCCAG GTTAGAGTGA
72421   AGTGTTGTGA TCATAGCTCT CTGCAACTTC AAACTGCTGG GCAAAAGTGA TCCTCCTGCC
72481   TCAGCCTCAT GAGTAGAGTA GCGGGAACTA CAGGCGCATG CCACTGCACC CAGCTAATCA
72541   CTATTTATTA TGCTCCTACT GTGTGCTTTA GTATATTTTC TGTTGTTTTC TGCAACCCAT
72601   TTTGAGGGCG TGTTAGGGAA TACAGATGCA GTAACTTTGG TCTCAGCCCT TGAGGTGAGG
72661   AAATATTTAG CCTCAGGTTT AATCTAATTG TTGGCCATTT GCCTTCAAAG ATTGAAATAT
72721   GAGCAAAACT GTGGCTCTGG GTTATATGTT AAAAAAAAGT TTATGGGGCT GAAGCCAGGC
72781   AACAGACAAG AGCCCCTACA ATCTTATTTA GGCTGAAAAT ATCCTGGAGT CCCTGTATTG
72841   TTGGTCTCAA GCAGATAGCA ACACTAACAC TTACTCTTTG AGGCAGGCAC TGCCAGTGGG
72901   GTGGCTGTTA TTATTAGCTT CATTAATTGG TGAGTCAGGA AAAAACAGCT TTAAATCATT
72961   CAAAGTTCTG GCCTATACAG GATTTAGTAA TATTAGGTTA GCTACATCCA AAAGATGACA
73021   GAACCCTACT CTAAGGCTGG GCTTGGTGGT TCACACCTAT AATCTCAAAA CTTTGGGAGG
73081   CTGAGGCAGG AGGATCACTT GGTGCCAAGA GTTTGAGACC AGCCTGAGCA ACATAGTGAG
73141   ACCCCTGTCT CTATCAAAAA CAAAGAACTC TAATTGGCAT AGTAGAAGGA AAAAGTGAAA
73201   GAAAAACCAG CTGTCACCCT CATTCCTTAC ACCTGTCCTA ACAACTCCTC TCACTATCCT
73261   TTGAATATAT CTTGGCTGTT TGAGTCTCTC TCTAGCCCCA TTACTGCTGT TTGGACTTGA
73321   CATTTTGCTC TGCATTTTTA ACTTTTCTAC CAGGGTTTCC AGACCCTGAA GAGTGTGGCA
73381   TGAAACAAAA CTAGTCAACC TATAATATTT ATGATGTGTG TGTAAATAAA AGAATACACA
73441   ATATATTGCA TTACAATATT TTAACTGTGT CCTCAATTTG TTTGTGGCTT TCTTGAGGAC
73501   ATCAGTTTTG GGTGGGACGA CCACATCCTT AATCTGAACT TTCCCTTGGA GGTCATTCTT
73561   TTTTTTTTGA AATAGAGTCT CGCTCTGTCA CCCAGGCTGG AGTGCAGTGG CGCAATCTCA
73621   GCTCACTGCA ACGTCCGCCT CCTGGGTTCA AGTGATTCTC CTGCCTCAGC CTTCCAAGTA
73681   GCTGGGATTA CAGATGCACG CCACCATGCC GAGCTAATTT TTGTATTTTT AGAAGAGACG
73741   GAATTTCACC ATGTTGGTCA GGCTGGTCTT AAACTCCTGA CCTCATGATC TGCCCACCTC
73801   AGCCTCCTAA AGTGCTGGGA TTACAGGCGT GAGCCACCCC GCCCGGCCAG AGGTCATTCT
73861   AATAGACTTT TTTTTGTTG TTGCTCACAG GCTTGTTCAA TCTTATTTCA AAATTTGAGA
73921   AATACAGTTT CCATGGAACA CCAACCAGAT ATCAGGTTGC TATGGAGTTG ATAGTCAAAA
73981   GCTTTGTATC TTCCAGTTTT TCAGAATGGC TTCTAAAGGT TCTGATTCAG AGCTCTTAGG
74041   CGAAATTGAA CAACCAAGTG TCAAAGTACA ACATTCAGGA AGTTAAAAAC ATGACTGACA
74101   TATATGTACT ATATATAGTG AGCTTGTGTA TGTGTCAATG AATGATTTAA TTCATTAATG
74161   AAGGAGGAAG CAGAATCACA ATTAGGTCAA AGGAAGATAC GGGAGAATAA AATATGTATT
74221   TGGTCAGGGA AAGGATGTAT ACTGGAAGAG GAAGGGAAAA TCAGATATAA AGTTGTTTAA
74281   TGACTTATTA GGCAATACAA TAATAACTTT TAGGGTCATT TTTTCTATAT TAAGAATTCA
74341   TTTCCATCTC TATGACAAAA TCCTTATTAA TTTATTAAAC TTCTACAAGT GAATGTTTAC
74401   TTTTAGATAG TCTGGACCCA ATAAAATGTA AACATTAAGT CAGAGTTACT TTCACGTAGG
74461   ACAGTGTTGT CCAATAAGGT ACCACTAGCT ACACGTGATC ATTGACCATT TGGACTATAG
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74521    CTAGACTGAT TTAAAATGTT CTAAAAGTGT AAAATACACA CCAGGTTCTG AAGATTTATC
74581    ATTTAAAAAA GAATGTCAAC TGTCTTTTTT TTTAGCTTAT TTATTATATG TTGAAGTGAT
74641    AATAGTTTAG ATATATTAAG TTAAATAAAA TATCTTAAAA TTAATTTTAC TTGTTTCTTT
74701    TCATTCTTTC AATGTGACCA CTAGAAATCT GGAAAGTATT TATGTGATTC ACATTCTATT
74761    TTACTGTCTA GTATTGCCTT ACATCATCAG GTACCCATA  AGTAGGCTTT TTAGATAATT
74821    CTCTAATATA GCTTGGAAGG ATATGGAGAA ATATTTTGC  GTTGCTTTTA AGTTTTGCAT
74881    AACTTTTTCA ACACACTTTA TAAAGGATCT AGAAAAGGGT TGGTTACATG TTTCTCTGTC
74941    TTCTGGCCTC CACCATGTTG CCAGGAGGTT GGGGACAAGA TTCTGGGTGG CTGGATGTCC
75001    TAATGGCTTG AGGTCTGGAC TTGAGATTTG CATATAAAGA GATGTGATTA GATTGAGTCG
75061    ACTAGAAAAA TCATATTAGA GAACTGAATC ACAGCGATTA AATTACATG  TCGATTTATA
75121    AACCAGGACA CCAATTTATA GTGAAAGAAG GTCCAGTTAC CTGGTAATCA AGACGTTTCA
75181    TAGCTATTTT CATGATGGAT ATACTTAGCT GAGTTTTAAA TGAGAAGGGG GTTCATTGCA
75241    CATAGAATAA GATCTAAGTG AAATGTTTAT TTATTTTTTT TTTTTTTTGA CATGGAGTCT
75301    TGCTCTGTTG CCCAGGCTGG AGTGCAATGA GGCAATCTCG GCTTCTGGAG TGCAATGAGG
75361    CAATCTCGGC TTCTGGAGTG CAACGAGGCA ATCTCGGCTC ACTGCAACCT CCACCTCCCG
75421    GGTTCAAATG ATTCTCCTGC CTCAGTTTCC TGAGTAGCTG GGATTAGAGT TGCCTGCCAC
75481    CACGCCAGGC TAATTTTTGT ATTTTTTTTA GTAGAGATGG GGTTTCACCA TGCTGGCCAG
75541    GCTGGTCTCG AACTCCTGAC CTCAGGCGAT CTGCCCGCCT CAGCCTCCCA AAGTGCTAGG
75601    ATTACAGGCG TGAGCCACCA AGCCTGGCCT AAGTGACATG TTCTTATATT GTTCCTTTCT
75661    TTCTTTTTTT TTCGACTGAG TCTCACCCTG TTGCACAGGC TGGAGTGCAG TGGCGTCATT
75721    TCGGCTCATT GCAACCTCTG CTTCCCGGGT TCAAGCGATT CCCTTGCCTC AGCCTCCTGA
75781    GTGCCACCAC CCCCAGCTAA TTTTTGTACT TTTAGTAGAG ATGGTGTTTC ACCATGTCCG
75841    CTAGGCTGAT CTCAAACTCC TGGCCTCAGG TGATCCGCCC CCGAGTCTCC CAAAGTGCTA
75901    GGATTACAGG CGTGGGCCAC GGGGCCCAGC CTTATATTAT TTCTTTTACT ACAATATATT
75961    AGTATGATGC AGGTGCTTCA ATTGTTTATA CACTTTCCAT AATTTTGTAT AATTCTTATA
76021    CCCTGTCACT CTGAGGAATA GCCGGTCTAA GTGTTTTTCC ACCACTGCTA ATTCATCCAT
76081    CACTAATCTC ATTAGACTGT TAATTCCCAG AGGACATAAG CACACAAGCA GACAATGTTT
76141    ACAAATGTTG GACAAATGTT ATTTAATAAA ACAATGGGGT CACCCTTAGT CTAAAAGATG
76201    TTTCACTTTT CATTTGTCAT TGAACTCTTA TTTGTAGGTT CCCTTTTGAC TTTCCCACAA
76261    TCTAAGGCTG TTCTCTTTAA CACATATTTT CATGAAAACA TATATTGAG  CAGAAATTGT
76321    TGGGGAGTTG TAATATTACC TTTGTCCCTA AATATGAATC TATAATTATA TCAAATATAT
76381    GGGCAGACAA TTTACTTTGC CTTTAATCTC AAGAAAAAAA TAGCAATTAC TTGGGGTCGG
76441    AGAGTAAAAT AAGAAGTAGT GAACCTTAAA GTAGCAAACT TTAGAACAGA ATAGTTTCAG
76501    AGGGGATGAG AAGAGGTGAT TTTTCAGCTC ATCAACAACA GATCTTATAA TAAATTACAT
76561    GTTCTGGTAC TTTTCTTGTC TTTCTGTGTT AAATTTTGCT ATTTAAAAAA ATAAATTTCA
76621    AATACATTGT TCATCTTAAA AGTCAAGAGT GTGTTTTATT AAAGTCAGTT GCTTTATTTG
76681    CAACTCAAAA GATATATTTG AGTTCCCAAC TGGAGATTGT CCTATATGGT AACTTGCGTA
76741    AGGTATGGTT ACTGAAAGTA ACCTACAATT TTCATGGGCT GAAATTCATT TCTATATTGC
76801    AGCGTACAAA AATAAATAAA TAAAAAATGC TTGTTTTCTT TGAAAACATA TTATCTCAGT
76861    GCCTCTAACT GCCAAATCTA TTGGCTTTTT TGCAGGCTTA AGGGCTCTCC CTTGTTCCTT
76921    TATGATCTCT ATCTTGAGGG CCAGACCTCC TGCCTTACAC AACTCAGAGG GGGACCTCAG
76981    AGCTCTTTAA AAAGAGCCCA ATTTCTCGCC TGTAGAGAAG TGAAAAGGAT GCCCCACCCC
77041    CATCTATGAA AAGAGGGATT TGATAGTTTC AATGTCTTCA AATCAAAGAT TTAAGTCTGT
77101    AGCCCCCAC  CACCCCGGAC CCTAGCAAGG CTCATGAACC CCCTCCCATC CCGCCCTAAT
77161    TGCTTTGGAC TGGCCGTGGA ATCCTTGTCC CAGTCCACAG TTCCTGTGCG ACTGCACGAA
77221    GAATTCACAG AGGACCTGTG TTACTTCCCT TGTGAAGAAA CAGAATTATC ATGAAAATTT
77281    AGGTGGAAAC CATTTCGCTT TTTTCTTCAA AAATAAGGGA AGCATGTGCC CAACCACCCC
77341    TGGGAAAAAG AACCTTCAGG GGCAAGGAG  CGAACAGGTA ATTTATAAGA AAAACAGAAA
77401    GTGGTCTCTG ACTGCCCCAG ACTTCCTTCG GAGTTGGGGG AATTGGGGAC GCCTGGACGC
77461    GTTGTTTTTG CGTTTGTGGA AAAAATAAAT GAAGAGCATG AAGCCCGAGG CTTCTGAGAT
77521    CCTTTCCTGA CCAAACCCAA GTGATTTGGT GCGGGGAATT TTAATATTTT TCCCCTTTTG
77581    TGAGGTGGAA CAAACACAAC TTGGGAGCAG CGCAGCGGCT CAGAGCCTGC CAGCCAGGCG
77641    GGCGACCAGA GCACCAATCA GAGCGCGCCT GCGCTCTATA TATACAGCGG CCCTGCCCAG
77701    ACGCTGCTTC ATCGGCGCTT TGCCACTTGT ACCCGAGTTT TTGATTCTCA ACATGTCCGA
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77761   GACTGCTCCT GCCGCTCCCG CTGCCGCGCC TCCTGCGGAG AAGGCCCCTG TAAAGAAGAA
77821   GGCGGCCAAA AAGGCTGGGG GTACGCCTCG TAAGGCGTCC GGTCCCCCGG TGTCAGAGCT
77881   CATCACCAAG GCTGTGGCCG CCTCTAAAGA GCGTAGCGGA GTTTCTCTGG CTGCTCTGAA
77941   AAAAGCGTTG GCTGCCGCCG GCTATGATGT GGAGAAAAAC AACAGCCGTA TCAAACTTGG
78001   TCTCAAGAGC CTGGTGAGCA AGGGCACTCT GGTGCAAACG AAAGGCACCG GTGCTTCTGG
78061   CTCCTTTAAA CTCAACAAGA AGGCAGCCTC CGGGGAAGCC AAGCCCAAGG TTAAAAAGGC
78121   GGGCGGAACC AAACCTAAGA AGCCAGTTGG GGCAGCCAAG AAGCCCAAGA AGGCGGCTGG
78181   CGGCGCAACT CCGAAGAAGA GCGCTAAGAA AACACCGAAG AAAGCGAAGA AGCCGGCCGC
78241   GGCCACTGTA ACCAAGAAAG TGGCTAAGAG CCCAAAGAAG GCCAAGGTTG CGAAGCCCAA
78301   GAAAGCTGCC AAAAGTGCTG CTAAGGCTGT GAAGCCGAAG GCCGCTAAGC CCAAGGTTGT
78361   CAAGCCTAAG AAGGCGGCGC CCAAGAAGAA ATAGGCGAAC GCCTACTTCT AAAACCCAAA
78421   AGGCTCTTTT CAGAGCCACC ACTGATCTCA ATAAAAGAGC TGGATAATTT CTTTACTATC
78481   TGCCTTTTCT TGTTCTGCCC TGTTACTTAA GGTTAGTCGT ATGGGAGTTA CTGAGGTATC
78541   AGAGACGAAT TGGGTGACGG GGTTGGAGAG TGGCCGTGGT GAGGTTACAG CATTTAAACC
78601   TTTATTGCGG CTTCTAGGTC CCTGACCGGA GGCTTTTCTC GCTGGCGGAT GGTTTTGGGA
78661   TGGCAGTCCC GCCCCAGGCC TGTGAACGGC AGAAAAGACC GCAAAACAAG AGCCAGTTTC
78721   TTAGTCTAAA GGGATGTCCG GATTGGACTA AAAAATTTTC AAAAGTCCCG CCCTGCTCCC
78781   GGGTTGGTCC GTTCTTCTAG TACATGACTT TCATTCTGTA TTTAATTGGA TGGTGGAAGA
78841   CGTTGCTTAT TCTGTGTTTT TTGCTTTACT GTGACTTAAA AGTTTGCCT CTTTTCTCTT
78901   TATATTAATG TCTGGGATTT CGGACGCTTT CCATGTTGTT GGTAGTCAAG TTGATGTCTC
78961   CTGGAGGTAG TGGCAACATC CAGCCCTGGG AGGAGAGTGC GTGCAGGTAC CTTTGTCCTA
79021   CATTCCTCTG CTGTTAATTT CTCATTCCTG TGGCAACGAA GGAATGCATT TAAAAAACAG
79081   CCACAACAGC GGCAATAGCC CTTCCTCCAC CCAAGGCAAT CGTGGACCTA GGGAGTTTTT
79141   TGTGCCACAT AACATGTAGC CTTCCGCTAA ACTGACAGGT TTGAGCGTAT CGATTTTGAG
79201   CGTATCGAAA GCACAACTTT TAGCCAGCCA TTTTGTCCTC GCATGACTAC GGTTGCTTAT
79261   CCTGTTTAGA CAGACAGCAA CATTTAAAAA TCGAAGTTCC TTTAAACGTA TTTTGTTTGG
79321   CAGTCCAAAT GTTTCTATGC AGAAAACAGT ATTTGTACTA TTAACTATGA AGAGTGTATG
79381   GATAAATGGG AGACATTTCT AATAAAGGCC TTCGTTAATG GTTCCCTCTG TTTGACATCC
79441   ATGGTGCTTC TGAATACAGA AAGCCTAGCG TCTTATATTC GCTTCTTTTA AAATCTGGTG
79501   GGCACATTTT GGTGAGACCT AAATTATGGG GACTGGGGCT TCTGGAGATA AGCTGCTCAA
79561   TTATTCTACC ATCTCCACAA TGATTAATAT AGTGAGTTGA TTTGTTAGTG ATAGTGACCA
79621   CGGATTCATC CCAAGAAAGA GAAAGGGGAG GGAGGCAAGC AGAGAGACAG GAAGACAGAG
79681   GCAGGGAAGA AGGAGAAAAC ATTCTCCCAT GGTTTAAGTA ATTTTGTGTT GTTAATTTTA
79741   CATTACAACA CGGTTTAACA TGGTGAACCC TCTATTTTGG TGTAAGGTTT AACATATGGA
79801   CATATTTTTC CCAAGACCAT TTATGAACTT TCATTTCTGC TTCCCCCTTC TTCCTCCCGT
79861   GCCACCCTCC ACGCTCCTAT CAATTTTGGC TGTTTTGTCA TAGGCTAATA CGCTATAATT
79921   TCATGGACAG TTGGACTGTC TTAGGTTTCT CAGGTTTCTA TTTTGTTCCT TTAGTCATTC
79981   CCACAATTCT TAAGGTAGAA TTGTATTGTT TTAAACATTG TGTTGTGTGC TATCCTCAAT
80041   GCTGAGATGA TTATGTGACA AATGGCAAGT GTTCAACTAA TACCTAAATC TGTAGTATCT
80101   TATCAAGCCT AATGCTACTT CACAATGCCT ACTCCATTCA CCGCACTTTA TCTCATTACT
80161   GGCATTCTGT CATCTCACAT CATCACAAGT AAAACGGTAA GCTATTTTGA GAGAGATCAC
80221   AGTCATATAA TTATATTTAT ATTTATTTAT TTATTTATGA GACGGAGTTT CCCTCTGTCA
80281   CCCAGGCTGG AGTGCTGTGG CACGTTCTCG GCTCACTGCA ACCTCCGCCT CACGGGTTCA
80341   AGCGATTCTC CTGCCTCCGC CTCCCGAGTA GCTGAGATTA CAGGGGCCTG CCACCATGCC
80401   CGGCTAATTT TTGTATTTTT AGTAGAGACG GGGTTTCACT AAGTTGGCCA GGCTGGTCTC
80461   GAACTCCTGA CCTCAGGTTA TCCGCCCACC TCATCCTGCC AAAGTGCTTA GATTACAGGC
80521   GTGAACCACC GTTCACAGAC TCAAATCATT TTTATTACAG TATATTGTTA TAATTGTTGT
80581   TTTATTATCA GTTATTGCTA ATCTCTTACA GTGCCTGATT TATAAATTAA ATTCATCATT
80641   GCCATGTGTA TATAGAAAAA AACAGTGTAT ATACGGTTCA GTACTATCTG TGGTTTCAGG
80701   CATCCACTGG GGGTGCAGTT TATTAAACAT GCATTACAT TAGTCTCCCC TTTGGGAGAC
80761   TAATTAACTG AGATGTTGTA ACGTGACTTT AATAGCAGAT AGAGCTAATT TTCTCTCATT
80821   ACTCTTCTTT TTCAGAATTT TCCTGGTTAT TCCATTTTTT ATTTTTCCAT ATGTATATTA
80881   AGATCTCTTC CACCTCCTCC TGTTTCTCCA TCTCAACATC AAACAATTAA AAAAAAAAAA
80941   AAAGGCTGGG CGCGGTGGCT CACGCCTATA ATCCCAGCTC TTTGGGAGGC CTAGGCGGGT
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81001   GGATCACGAG GTCAGGAGTT CAAGACCAGC CTCGCCAAGA TGGTGAAATC CCGTCTCTAC
81061   TAAAAGTATA AAAATTAGCC AACCATGGTG GCAGGCGCCT GTAATCCCGG CTACTCGGGA
81121   GGCTGAGGCA GAGAATTGCT TGAACCCGGG AGGCGGAGGT TGCAGTGAGG CGAGACCTTG
81181   CACTCCAGCC TGGGTGACAC AGCGAGACTC CGTCATAAAA AAAAAAGCCG GAAGCAGTGG
81241   CTCACGCCTG TAATTCCAGC ACTTTGGGAG GCTGAGTCAG GCAGATTACC TGAGGTCAGG
81301   AGTTCAGGAC CAGCCTGGCC ATGAAAATAC AGCCTGGCCA TGAAAACACA CAATAAATTA
81361   GCTGGGCGTG GTGTCACACA CCTGTAATCC TAGCTACTCG GGAGGCTGAG ACAGGAGAAT
81421   CACTTGAACC CAGGAGGCAG AGGTTGCAGT GAGTTAAGAT GACGCCACTG CACTCCATCT
81481   GGGCGACAGA GCCAGACTCT CTCTCAAAAA ACTAAATAAA TAAAAATAAA GTTATGGTAC
81541   ATTGAACTTC TGTGTTCCTT TCTCCCTTAG ATACTTTCAT GGCTACCCAT TTAATTGATG
81601   TTCTTATCAT CTCCAAGAGT TAGTCAGGAG AGGAATCAAC CCAAGCAAAA ATAGCTGATT
81661   TTCTAATTTT CCTTCAATGC CCTTTGGGGT CTTAATCCAT TTGATTTATG TACTTTCAAT
81721   TAATCCTAAC CTCGAATGTC TTCTGCAAAC ATGTTTCCAC AGATGAAACT CGTCAAATGA
81781   AACACATTCC TTTAATTTAT AGAGTTAAAA ATTAGAAAAA TTTTCAATTC TATTTGGCCT
81841   TTAGATTCAG TCTTGCATAT GTTTTCTCAA TTTTGTTCAT GCTCTTTAGT TTTGTTTTAT
81901   TCCATCACAA TTGTTCACAT AGCTTACTGG CTTAGGTCTA ATGAACCATT CATTTGGAAA
81961   TTAAAATTGG CCATTTTAAG ATGAAAAGA TTCTTGCCTC AATTTTACTT AGTTTTTGAA
82021   ACTGTCAATG AGGACACATG TTTTTCTGTA CTCTTAGATT CACTAAGTAG TGTCTTGCAA
82081   ATTTAACTGA CAAAGGACAG ATTAACATGC GAAAAAAAAA GCATGCAATT TTATTAGTAT
82141   ATTACATGCA CAGAGTTCCC AAAGAAAAAA AAATTGAAAC CTTAAAAACG CGGTTAGACT
82201   CACAGACTTA TACACCATTC CAACAAAGGA AAGGGAGTTT GCACTTCATG GGATGACGAA
82261   TTTGGGAATG TGACAAGGAA ATAAATACAT GGGCAATAAA AACCATGGAA GATAAAATGA
82321   AAGATAGAAA TAATTGTAGT AAGGTTTGTT TTTGCAGAGT CATCTCAGTG CCAACCTTCC
82381   ATATCTAGTG ATAAGAATTG CTCTCTTTTT CCTGGTATAG CAGTTGGGGA CACTTTTACA
82441   AGGGAAATTT CTGTCACCTT CACAAAGGGA AATTTGGGTA AAGAGAAGAC AGAGACCTCT
82501   TCCTACACCT GTTGATTTTC AATTGCCTTC AGCTGAAAAT AACTTTTATG CCAAAGTAGA
82561   ATAATTTGGG GGTGACATCC TGATATTCTT CAAAACTTAT ATTTAATTTC ACATTAGTAA
82621   TTATATCATT TTTGATTTTT AAATTAGTTT TATAAAATAA TTTTGAAAAA CGGTAATAAT
82681   ATTCAAATAA TTCCAGAAAC ACTGCTGATA AGCCAAAAAC ATCAATGAAT ATTGCATAAA
82741   CAACTGATAA TTCAACCATG AAAATTTATG ACATTGTTCT TGTGTGATAA AACTATGAGT
82801   AACATAAAAA CTAGAGGCTA CTTGTAATGC ATTATTCCAA ACTTTCTGTT TTTTATTTAT
82861   TTATTTATTT ATTTTGAGAC ATAGTCTCTC TCTGTCACCC AGGTTGGAGT GCAATGGCGT
82921   GATCTTGGTT CACTGCAGCC TCCACTTCCC CGGTTCAAGC AATTCTCCTG CCTCAGCCTC
82981   CTGAGTAACT GGGATTACAG GCACCTGACA CCAAACCCGG CTAATTTTTT TGTATTTTTA
83041   GTAGAGACGG GGTTTCGCCA TGTTTGCCAG GCTAGTCTCG AACTCCTGAC CTCAGTGATC
83101   CACCTACCTC GGCCTCCCAA AGTGCTAGGA TTACAGGCGT GAGCCACCAT GCCCGGCGCA
83161   TTATTCCAAA CTTTCATACA CAGTGCTATC ATGGCTACAA ATTGAAGTAT CATATTATAC
83221   ACTCCTAGGC AAAGCTCTGG ATATTTGGC TATATAAGCC TGAGGGAAAT GTAGTAAGGA
83281   CATTGTGGTT GAAATTCATA CCAGAGATGA ACAGGCCCAG TGCAAGACAG AATTACATCA
83341   CTAAAGGATA TCAGAAGAGA ATAGGGATTT AGGGTACAGT GGCAACAACA GTTTTGGGAA
83401   CTAGCATTTT TTGAGCACTT ATTTACAATA TGCCAAGCAC TGTTGCTGAT TACTCTATAT
83461   TTATTTTCAA ACACATTCTT GTCACAGCAC TTTGAAGTAA GTGCCATTGT CATTCCCACT
83521   TCAGGGTGAA GGACTAAAGC TTGGTGTCAT TAAGGATGTA GCTAGTTAGC TGTGTGTGTG
83581   TGTGTGTGTG TGTGTGCATT TTTTTTTAAA TTTAAAGTCA ATAAATTTTT ATTTGAAGAA
83641   TTTCACATCA AGGTAAACTT TGTTCCTCTA AAGAGCTGGA GTCAAAATGT ATCTTCAAAA
83701   GATTCATCTT CAAGTTAGCC CTTCTTAATA GAACTGATGC TTAATCCACA GTTGTCAGCC
83761   CACAGTTCTT TTATTTTGAC TTTTTTTTTT TTTTTTTTTG AGACGGAGTC TCTCACTGTC
83821   ACCCAGGCTG CTGGGCAGTG GCGTGATCTC GGCTCGCTGC AACCTCTGCC TCCCGGGTTC
83881   AAGTGATTCT CCTGCCTCAG CCTCCTTAGT AGCTGGGACC ACAGGCGCAT GCCATCGTGC
83941   TCGGCTAATT TTTGTATTTT TATTAGAGAC AGGGTTTCAC TATGTTGGCC AGGCTGATCT
84001   CAAACTCCTG ACCTCATGAT CCGCCTGCCT TGGCCTCTCA AGTGCTGGG ATTACAGGTG
84061   TGAGCCACTG CACCCGGCCT TATTTTGCCT TCTTTAATCT CCATTTGAAC ATGGACATAC
84121   TGATGAAAAC TACAACATTC TTCACCAAAA ATCTTTGGGA TTTAATTTCT TCAACCACTT
84181   TACTTTGGGG TCATTTTAAG ATTAGGTGTA TCTGCCTGGT TCTCAATTTG ACACCCTTTC
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84241   TCTCTAAACA TGAATGAGTT CCAATCATAT TTATTCCTAA GCTATCACAC TCAAATATAC
84301   TACAGATCTG TGGAATATGC CAAAAGTTAA GGTGAAAAAT TAAATTATTA GGTATTTCAT
84361   AGTTTTGCTA GTTTTTGATC TGTGAGTGAA TATAACTATC CTCTATGTCC TGGCACTGTT
84421   CCTCAGAAAC ATAGGGTCCA CATATGTAAT TTTAAATTTT TTAATAGGCA CATTTTAAAA
84481   AGTGGAAAAA GAAATCTATT TTAATGATTT GAATCCAGTG TAACCAAAAA TTGTTTCAAC
84541   AAGGTATCTA ATATTAAAAT ATTGAGTTTT TACTTTGTTA TTTTACTAGG TCTTTGAAAT
84601   CTGGTGTGTA TTTTACACTT AAAGCACATC ACAGTTTGGA GTAGCCACAT TTCCAATGCT
84661   TAATACTCAC ATATGGTTAG TGGCAACTAT CTTGGACAGG ACAGCTTTTA TACTCTGGGA
84721   AGACACAAGC AAATACTTGC TCTGCAGCAG AATCCAGATG TTTTCCAAGA AAACACTTTT
84781   TCTGACCTGT TCGTGAAACC CAGGTAGTGT CTCTAATACT TTATATTTTA TTGGTTTGTC
84841   CTATTGTAAC CACCCAACGG GCTCTCCTTG TCCACTTCCT AGACAGAGCT GATTTATCAA
84901   GACAGGGGAA TTGCAATAAG GAGCCAGCGC TACAGGAGAC TAGAGTTTTA TTATTACTCA
84961   AATCAGTCTC CTTGAGAATT TGGGGACCAA AGTTTTTAAG GATAATTTGA TTGTAGGGGA
85021   CCAGTGAGTC GGGAGTGCTG CTTGGTTGGG TCAGAGATGA AATTATAGGG AGCCTAAGCT
85081   GTCCTCTTGT GCTAAATCAG TTCCTGGGAG TGGTGGGGTG GGGGACTCAA GACCAGATAA
85141   TCCAGTTTAT CTATATGGGT GGTGCCAGCT AATCCATTGT GTTCAGGGTC TGCAAAATAG
85201   CTCAAGCATT GATCTTAGGT TTTAAAATAG TGATTTTATC CCCAGGAGCA ATTTGAGGTT
85261   TAGAATCTTG TAGCTTCCAG CTGCATGACT CCTAAACCAT AATTTATAAT CTTGTGGCTA
85321   ATTTGTTAGT CCTGCAAAAG CAGTCTGGTC CCCAGGCAGG AAAGGGGTTT GTTTCTGAAA
85381   GGGCTGTTAT TGTTTTTGTT TAAAAGCAAA AGTATAAACT AAGCTCCTCC CAAAGTTAGT
85441   TAATCCCAAA CTCAGGAATG AAAAGGACAG CTTGGAGGTT AGACGTTAGA TGGAGTCGGT
85501   TAGGTAAGAT CTCTTTCACT GTAATAATTT TCTCAGTTAT GATTTTTGCA AAGGCAGTTT
85561   CACTGTCCAC TTCACCTCAC ATCAGGCCTC TGACTAGAGG ATTCCAACAA TACTTAGGCC
85621   AGGACACCAC CATGTCTCCT TATCCACCCT GAGGGATTCC AATTTCTGAA ACAAAGGAAA
85681   CTATATATGA TAGTATGAAA CTATATATGA GAAGGAAATT ATATATGATA ATCAATTTTA
85741   GGGTTATCTT ATTGATTAGA AGATATTAAA GTGTGACACT GCCTGGCAAT GATATCTGCT
85801   GGTAGTAAGA ATTTGGCGAA TTTAGTGAAA TTCCTGAGGC TGAACCTCCA CTTCTGTAAA
85861   ATGGAGACAG TGAGATAATT TGCCTTACAA TGCTGAAGTA AGAATTTTAC ACAATAATTC
85921   AGACCAACCA CTTCATGTGG TACTTGGCCC GTGGAAGACT ATCAATGACA GTTAGTTTAT
85981   AGTTTATACT ATTAATGAAT CCTTTGTTTC ATTGTTATTT CCTTCTACAC GTTGGCCTCT
86041   CTAAAGAAG GTAATATTCA ATACAAATAA AGTTAAAACA GCTTGCAGAG TTGTCCCAGG
86101   GAACTCACTT AACCACTGAA GTGTTCAAAT TGCTTAAGGT TGACTTTATA TTCTCCTGAC
86161   TAACCTTTCT CCTTCTGGTA TTTCTTCTGA GAACAGCACC ACCATCCAAA GCATCATGCA
86221   AACAGTGGTC ATCCCAGACC AGTAATTCTC AACTCACAGG GTGCTCCTGC AGAGATGTAT
86281   TTGAATAGAG TGGTAGGATG CTGAAGAAGG CCACGTAAAA TTTGGCCAGT GATCTGGGGC
86341   AGATTTATCC TGAAGCTAAT GAAACACAAG TGTAAGGGCC TGTACTTCCA AGGTGCAGAG
86401   AGGGGCCCTA CAAATGTGTT AGTTTGTCTC TCTCTCTCTC TCTGATTTTA AAATTTGCAG
86461   TATTAAGGTA CTTTAATCAC GGATGGTTCA GGCTGCTATT TTCACTCAAT CCTCCTTTTT
86521   ATTAAAATCA CCATTGTCTG ATTATGTTAG AATCCTGATG AAAATATTTG GAATTTGAGT
86581   AAGAGAAAGT TTAGTTGAAG ATGTATCTAG TATGGGGATA ATAAGTTACG TGATTTGCAT
86641   ATGTGATCAT GTGTACTTCA TTCGTTGCCA GCCAATCTGA CGTAAGAATG GCTTCAAGGA
86701   GGCCGGGCGC GGTGGCTCAC GCCTGTAATC CTAGCACTTT GGGAGGCCGA GACGGGCGGA
86761   TCACGAGGTC AGGAGATCGA GACCATCTTG GCTAACACGG TGAAACCCCG TTTCTACTAA
86821   AAATACAAAA AATTAGCCGG GCGTGTTGGC GGGCGCCTGT AGTCCCAGCT ACTTGGGAGG
86881   CTGAGGCAGG AGAATGGCAT GAACCTGGGA GGCGGAGCTT GCAGTGAGCC GAGATCGCGC
86941   CACTGCACTC CAACCTGGGA GACACAGCGA GACTCCGTCT CAAAAAAAAA AAAAAAGAA
87001   TGGCTTCAAG GAATGTTCCT ACTGCTCACT GGAATAACTC ACCTAAATTC CTGGCAAGAT
87061   GCAGGTCTAG ATAAAATGTT ATGACATCTA AGTATTCAAA ACACATTCCC AGCACTGAGA
87121   GTGAGTGTCT AGTGGAGAGT AGAAACGTAT AGAGCCAGAA GCTAGTCTGG AAAGAATTCT
87181   TACAAAGTTT ACAACTTACA TGTGAAAGGA GCTTAACAGA GGATTTTCCA AATTTGAAAA
87241   CAATCCTAAA AACTTACTTG ACATTACCAA TAATGTGTTT TGAAACTGAA ATACTTCTAA
87301   GTTATGAAGA AAACATATTA TCATCAGCCA CCCTGGAGGA AAGATTGAAT TCTATTTCCA
87361   TTACCTATAG ACAACATTAC AAAATAATTT CGATCTGAAG ATGGAATCAG AGTATTCAGT
87421   CAAAACTACA GGAAAATATA CTTGGTAGTG TCATATTCAG AAGTTAATAA AATATGCTAT
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87481  TTTCTGAATT  TTGTGATGGC  TGTTGTTTTG  TCAGCTTTTA  TAAAATTGGA  ATTTGATTTT
87541  ATTTTCCCAT  TATAAATTTA  TATTTACAGT  CTGCAGTACT  TTTGCATTTT  TAATTTTACA
87601  TTATAGCTTT  TAATAGTTAA  CAAGTTGTAA  AAGGTTTGAT  CCCCAGAAAA  CCTTGATCTA
87661  CCCCCTCAGT  TAAGTATACT  AATATATTTA  GAAAATGGAT  GAAATCAGCA  TTTGAATATT
87721  TTTAAATATT  TATTAAAAGA  GGACATGGGT  AAAAGAGCTT  TGCAGTTGCC  ACCCTTCATT
87781  CTCAAATTCC  CTGGATAAGG  ATGACCGCAT  AATCTTTGGA  TGGTCATACG  CAAGTCTTGT
87841  GTATTTGTTA  CATAAATCTA  TTTAGTGGAC  TTTTGGCAGT  GTGTACTGAG  GCCAGTTTCT
87901  TCCACCTGAG  CTCTGACTCC  ACCTCCAGCA  GCCCAAACC   AATACTGAAT  TTTGGGGTCA
87961  GCTATTGTTT  TTGTGGACTT  AGGTAACTAC  ACACACATTG  TCTTTATGAT  AGCTTTAATA
88021  ATACTGCCAT  CAGAACTAAA  ATTGTCACGT  GGATTAAAAG  GAGTGACGGT  GGTGTCCCCA
88081  GGAGCCTTTC  AATATGTAAG  TATTTACACA  TATACATGCT  AAAAAGACCC  CTAGGAATTT
88141  TTTTAACAAG  GGCAAACAG   TAACTCAGCT  TGTTTTCTCG  CAGTAAAACC  GGTTGAAAAG
88201  GCCTGATAGA  CTTGTCTGCA  GTTACAAAAC  TTGTGTGTAG  TTATCACCTT  TATATCTCCT
88261  GGAAACTAAC  ATAGACAACC  GAATGGGTTA  CAACTGTTTT  TAAGTGAAAT  TGTGAGTGGC
88321  TCTGAAAAGA  GCCTTTTCAA  TGAGGAAGAA  ACGGGCAGAC  TTATGCCCTT  TCCCCACGGA
88381  TGCGACGTGC  CAGCTGGATA  TCTTTGGGCA  TGATGGTGAC  GCGTTTAGCG  TGAATAGCGC
88441  ACAGATTGGT  GTCTTCGAAG  AGTCCCACCA  GGTAGGCCTC  GCAAGCCTCC  TGCAGCGCCA
88501  TCACCGCAGA  GCTCTGGAAA  CGCAGGTCGG  TTTTGAAGTC  CTGGGCGATT  TCTCGCACCA
88561  GGCGCTGGAA  CGGCAGCTTC  CGGATCAGCA  GCTCGGTGGA  CTTCTGGTAG  CGACGGATTT
88621  CGCGCAAGGC  CACGGTGCCC  GGGCGGTAGC  GATGAGGTTT  CTTCACGCCA  CCGGTGGCCG
88681  GAGCGCTCTT  ACGGGCTGCT  TTAGTAGCAA  GCTGCTTGCG  CGGAGCTTTG  CCGCCGGTAG
88741  ACTTGCGAGC  TGTTTGCTTC  GTACGAGCCA  TTTGCAATGA  GAGCACACAC  AAAAGTGTAG
88801  TGAACTGAGA  GCAAGTGGCC  TTTAAATATA  GTGAGAAACA  TTCTGATTGG  TCCTGTAATA
88861  TTTCAAAAGT  CCCGCGCGAT  AAAATCATTG  GCTGAAGAGT  GACCAGACTG  ATTGGTTCAT
88921  TACTAGACAA  TCTTATTGGA  TGAGTTGCCC  CACCGCCCAT  CCTGTCCTTT  TCGTTTCAGT
88981  TATCTGCAGC  GACAAATTGT  CTAAAATTCT  AGTTCATCCA  GTCCCAAAGA  ACAGAGTGTA
89041  TAACAAGGTA  TCTAAGGATT  TTTAAAATGT  AAATTCCGAT  TCAGTAAGTT  TGAGTGGGAC
89101  TTGAAATTCT  GCATTCCTGA  CAGTCTCGCA  AGTTATCAAT  GCTGGTGAAC  ACTCACTAAA
89161  CCACCAGAAA  CGTTCAGACT  CATGTCGGGA  AATAACGCTT  ATATTCAGAG  AATGAGATTC
89221  CATGCTATTT  TGTTACTGGC  GAACAGCAAG  TTTCCTTGCC  CTTTGTTTTC  TAAGTCCAAG
89281  TCACATTCCC  ACCCTGCCTG  TTCTCAAAAT  GTCTTATTTT  GGTTGGCCTT  AAGTTTCACT
89341  TTGTATACTC  TAAAATGTAC  TTTCTAAAGG  AAGGTGTTAT  TTTCTCGAAA  CTTAACTTTT
89401  TAACACCATT  AGGCTAGGGG  GGCGGTGGCT  CACGCCTGTA  ATCCCAGCAT  TTTGGGAGGG
89461  CGAGATGGGA  CGATCACTAG  AGGCCAGGAG  TTCAAGACAA  CCCTGGCTAA  AATGGTGAAA
89521  CCCCGTCTCG  CATAAAAATA  CAAAAACTAG  CTGGGCGCGG  TAGCAGACGC  CTGTAATCCC
89581  AAGTACACAG  GAGGCTGTGG  CATGAGAACC  GCGTGAAGCG  GCGGGGTGGA  GGTTGCAGTA
89641  AGCCGATATC  GCGCCGCTGC  ACTCCAGCCT  GGGTGACAGA  GCTAGACTGT  CTCAAAACAA
89701  ACCAATCCAA  ACGAAAAGCA  AAAAATACCC  TAACAGAAGC  AAGTTATCAT  CCTTTCTTGT
89761  GTAACTATGG  ACGGCTCTGA  AAAATGCCGT  TTCAAGTGTA  AGCTACGTTT  TCTGATTTGA
89821  GTGTTTACTT  GACCTTGGCC  TTATCGTGGC  TCTGTTATTT  TGGCAACAGG  ACGGCCTGAA
89881  TATTGGACAG  GACGCCTCCC  TGAGCAATAG  TGACGTTGCC  CAGCTGCTTG  TTGACCTCCT
89941  CGTCGTTTCG  GATGGCCAGC  TGCAGGTGGC  GGGGGATGAT  GCTGCGGGTC  TTGTCACGTA
90001  TGGCGCTGCC  CACCAGTTCT  AAGATCTCGG  CGGCCAGGTA  CTGTAAGTAC  ACTGGCGCAC
90061  CGGCTCCGAC  CGGCTCAAAA  TAATTGCCCT  TTCGAAAAAG  ATGACGGACT  CTGCCCTATT
90121  GGGAACTGCA  AGCCCGGTAG  CGACGAACAA  GTTTTTGCTT  TAGCTCCATT  TTCCACGTCC
90181  GCAAATAGCG  ACCTATGAAA  GCAGCGGAAA  ACTGTGAAAG  ACAAGCAAGC  TGGAATGGCG
90241  CCTGAACAAA  TCCTTTTATA  CAAACTGCAA  GGCTGCAATA  GGAAGCTATC  CTATTGGTCA
90301  ATTATGTTTG  GTGCTTTATC  CAATAGAAAA  AGATAACATA  AATTCCATAT  TTGCATAAAC
90361  CCCACCCCTC  AGTGAAACCG  TGTTTCTTTT  GTCCAATCAG  AAGTGAGGAA  TCTTAAACCG
90421  TCATTTGAAT  CTCAGGACTA  TAAATACATG  GGCTCTGAAC  TGTTCTCTGT  ACTACTCTGT
90481  AGTGGAGAGT  GTTAGTAGCT  TTTCTATTCT  GTTTAGGAAT  AGCAATGCCT  GAACCCTCTA
90541  AGTCTGCTCC  AGCCCCTAAA  AAGGGTTCTA  AGAAGGCTAT  CACTAAGGCG  CAGAAGAAGG
90601  ATGGTAAGAA  GCGTAAGCGC  AGCCGCAAGG  AGAGCTATTC  TATCTATGTG  TACAAGGTTC
90661  TGAAGCAGGT  CCACCCCGAC  ACCGGCATCT  CATCCAAGGC  CATGGGGATC  ATGAATTCCT
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90721   TCGTCAACGA CATCTTCGAG CGCATCGCGG GCGAGGCTTC TCGCCTGGCT CACTACAATA
90781   AGCGCTCGAC CATCACCTCC AGGGAGATTC AGACGGCTGT GCGCCTGCTG CTGCCTGGGG
90841   AGCTGGCTAA GCATGCTGTG TCCGAGGGCA CTAAGGCAGT TACCAAGTAC ACTAGCTCTA
90901   AATAAGTGCT TATGTAAGCA CTTCCAAACC CAAAGGCTCT TTTCAGAGCC ACCTACTTTG
90961   TCACAAGGAG AGCTATAACC ACAATTCTT AAGGTGGTGC TGCTGCTATT CTGTTTCAGT
91021   TCTAGAGGAT CAACTGGAAT GTTAGCGAAG ACAAGTTTTA GAGCCAAGGT TAACTTGGAC
91081   GGGGCCGTGC GCGGTGCCTC TTGCCTTTAA TCCCGGCAAT TTGGGAGGCC GAGGCGGGCG
91141   GATCACTTGA GGTCGGGAGT TCGAGACTAG CCCGGCCAAC ATGGCGAAAG CCCGTCTCTA
91201   CTAAAATACA AATGATAGAC GGTCGTGATG GCGCTCTTTC TCATCTGTCT TAGCAAACTT
91261   CTTTGTTCCC CCTGGGTAAG CCTTCGGGTA CTATGTATAA TTCCTTTGAT AAGGTCACTA
91321   CTCCCTCCCT GGTCTAGTAC AGGAAACTTC CCTTTCTGGA TAATGAAGCA GGTAATGGAA
91381   TTCAGGGTAT AGTGTTCCTG TGGGGGTCAT TAGCCGTTAA CTTCTTGTGA GATGCGGGGG
91441   AGGGGAGCAG AAAAGTCTAA GCGACAAAAG GGCATGTAGG GATATTTGCT CCTGCAGCTT
91501   GCCTATGCTG TAAATTCTTA CTTCAAGTAT TGAGGAAACA ATAAGCGAAG TCTGATTTCC
91561   CGGGCGCCTT TATACGGAAT ATTTCCCGCT CCACAAAATG AAATCGCAGT AGTTTTGAGT
91621   TATAATTGTT TATCAATGAC AACAGCTATG TAGTTTACAT ATTTCATGCA TCCCAGAAAT
91681   CCAGATTCCC ATTTCCTAAG CCACTTAACG TTCTGATTTC CAGCTCTGCG AGATACAAAA
91741   GGGTTTGGAT TTTGTGCCCT TCCCCATCTG GCGCCACTGC AAAGCTTACT AGGAGGGCCC
91801   CACTTGGAGA GGGAAATCTT TTTCGAGAAG TCCAGGACGC CAAAAACAAT ATAGCTAAAA
91861   AAAAAAAAAA AAAAAAGGCA GGAAGAGCAC TAGTTGAGGA GGAGGACTCA ATGGGCCAAT
91921   TCTGGGGCTG GGGCTGGGGG AAGAAATGCA AGAAGAAAG ACACTTGTTG ACTGCACAGT
91981   AAGCAGGAGG GGGTGGGGGA ATCGGAGGGG AGTATTTTCA GCGAATTTAT GGGCATTATA
92041   TGTAGGTGAC ATACAGCAGT GTCTTTGGAT GAAGAAATAA AGTTTCTCAA ACAGTTCTTG
92101   TTTTTGTTTT GAGAAAGGGC CTTTCTCTGT CGGCCAGGCG CCATCATAGC TCACTGCAAC
92161   CTCGACTTCC CCAGCTCAAG CGATCCTCTT ACTTCAGCCC CTTGAGTGGC TGGGACTAGA
92221   GAAATGCACC ACCATACCCA GTTAATTTTT TAATTTTTTG TGGAGGCAAA GGGTCTTACT
92281   TTGTTGCCCA GGCTGGTCAA GCGAACTCCT GGGCTCAAAT GATCCTCCCG CCTTGGCCTC
92341   CCAAAGTCCT GGGATTATAG GAATGAGTCA CCGCGCCCGG CCCAGATTTA ATTTTTAAGA
92401   ATCTTTTAAA AGAGGTTCTG GGCCGGGTGT GGTGCAGCTC ACGCCTGTAA TACCAGCATT
92461   TTGGGAGGCC AAGGTGGGAG GATCACTTGA GCCCAGGAGC TCAAGACCAG TCTGGGCAAC
92521   TTAGTGAGAC CTTTTGTCTC CACCAAAAAT TTAAAAAATT AACCAGGCCT GGTGGCACAT
92581   TTCTGTAGTC CCAAGTACTG GGGAGGCTGA AGTGGGAGGA TCATTTGAGC CTGGAAGGTG
92641   GAGGTTGCAG TAAGCTGTGA CGGCACAACT GCACTCCAGT CTGGGTGAGG ACAGACCCTG
92701   TCTCAAAAAT AAAAAATAAA AAAAAATCTG GATGCCACAC AAAATGTCAG TGAACAACTG
92761   TAAGTGAAGC ACTTCCCATC CTAGTACTGT ATATGCAAAC TGCCGTTGTG AAAGTGACGC
92821   TTGGCTTAAA AATCTACATT CTTTTTTTAA TTATAAAACT ACCACATCCC CCAAAAACAT
92881   TACTAAGGAA TTGAGGCTGC AGTTTAAGAA GCTGATATTT AGGATCTATC TCCGGAGAAG
92941   TGAGACCTGG TAATATAAGC ATTTTCAAAA TGAACTTTTG GGCCAGGTGA GGTGTGTCAT
93001   GCCTGTAATC CCAGCACTTT GGGAGACCTA GTCAGGCAGA TCACTTGAGC TCACAATTCG
93061   AGACCAGCCT GAGCAACATG GCGAAATCCA GTCTCTACAA AAAATTAGCA GGGCGTGGTG
93121   GCATATGCCT ATAGTTCCAG CTACTATAGA GGCTGAGGTG GGAGGATTAC TTGAGCCCGG
93181   GAGGCAGAGG TTGCAGCAAG CCAAGATCGC GCCGCCACAG CCTGAGCGAC AGAATGAGAT
93241   ATGCACCCAC GCCCTAAAAA AAAGCATGAC TCATTAAAAA AAAAAAATTT AGCCGGTCGC
93301   GGTGGCTCAC GCCTGTAATC CCAGCACTTT GGGAGGCCGA GGCGGGCGGA TCACGAGGTC
93361   AGGAGATGGA GACCATCCTG CTTAACACGA TGAAACCCCG TCTCTACTAA AAATACAAAA
93421   TAATTAGCTG GGCGTGATGG TGGGCGCCTG TAGTCCCAGC TACTCGGGAG GCTGAGGCAG
93481   GAGAATGGCG TGAACGCGGG AGGCGGAGCT TGCAGTGAGC CGAGATCGCG CCACGGCACT
93541   CCAGCCTGGG TGACAGAGCG AGACTCCGTC TCAAAAAAAA AAAAAAAAA AAAATTAAAA
93601   AAATATGAAG TTTTGAAGCA GAAATTATTT TGTCGTATGT TCTTTCATAA ATTTTTTGCC
93661   TGCCTGCCTT CTTCCTTTGT TACAGAACTC CAACACTTAC CCAAAGGTAG CTGTTGGGTC
93721   AGGGTTTCTG TACTATAGTC CCTTCTGTGG TGGCCAGAAA TATGTTACAG GAAAGAGGTC
93781   CCCATCCAGA CCCCAAGAGA GGGTTCTTGG ATCCCGCGCA AGAAAGAGTT CAGGGTGAGT
93841   CCGCAGTGCA AAGTAAATGC AAGTTTACTA AGAAAGTAAA GTGGTGAAAC GACAACTACT
93901   CCATAGACAG AGCAGGACAT TCCCGAAAGT AAGAGGAGGA AGGCATCCAC CCTAGGTACA
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93961  ATACTTGTAT ATATGGGGAG ATGTGCTCTG CTACAAGTTT GTGATAAAGG ATTAATTTTC
94021  TTAGTTACTA TATTTTGCAA GAATCAACAT TATTATCTTT AAACAAAATT AAGAATGCCT
94081  TTGTTCTCCA GATATAGGGA TATCTGGACA CTCCTAAGTC TGAGTCTGTT TAGTAAACAT
94141  TATTTATTTG TTCCCTTAAC CGTAAACATC TAGAAGCTAG GAATGACTGA CTTTCTGGGA
94201  ATGCAGCCCA GAAAGTCTCA GCCTCATTTT CCTAGCCCTC ACTCAAAATG GAGTTACTCT
94261  GGTTCAAGTA ACTCTGACAC TTTTCTTCTC TTTTTTTCTT CTTTTTTCCT TCCTTTATTT
94321  TTTATTTTTT ATTTTTGAAA TAAGAAATCA AGAATACTTG ATGTTTCATC TAAAACAATA
94381  CCCATAATTG ATAAGCCAAA ACAAAAACCT AGGTCTTCTA ACTCAAAACT AGGATGTTTT
94441  GCTGTCTCTG CTGATACTCG GCTGATCGTT AATAGGTAAT TAACAAACAA GCCTTGCTAT
94501  GTCCCCCTCA GTTTATTACC ATTAGATCAT ATGCCTACTG TCAATCATAT TAATCCACAA
94561  CTATGCATTT CACAAAACTT GCCATAAAAA TTCACAGGTT TCCCGCTTCC CTCGAGTTTT
94621  CATTTCCGAA GGGTCCCATG TAATATAAAA CTTATATTAA ATACATTTGT ATGCTTTTCT
94681  CTTGCTAATC TTTTTTTTTG TTTTTTGAGA CTGAGCCTTG CTCTGTCACC CAGGCTGGAG
94741  TGCAATGGCG CGATCTCGGC TCACTGCAAC CTCCGCTTCC CAGGTTCAAG CGATTCTACT
94801  GCCTCGCCCT CCCGAGTAGC TGGGACCACA GATACGTGCC ACCATGCCCC GCTAATTTTT
94861  GTATTTTTAG TAGAGACAGG GTTTCACCGT GTTGGCCAGG ATGTTCTCAA TCTCCTTACC
94921  TCGTGATCCG CCCGCCTCGT CCTGCCAAAG TGCTCGGATT ACAGACGTGA GCCACTGCAC
94981  CCGACCAATC TGTCTTTTTG TAGAGGGGCC TCAAGCATGA ACTTACTGAT GGGTGAGAAA
95041  AACAGAATTT TCTTTTCCCC TACAATATAA ACATTAATTG TAATGTTATC ATTCAGGACA
95101  TTTTGGTGAC CAATCTTACA GAAATTTTAT CTTGTGCAAG TCTATGCAAA CCAATATGTA
95161  AATCTTCTAT AAGTGAGATT GTATTCACT TTTCTAGTAT CCTTTTAAAT TAATAAAAGA
95221  GATTCTAATG ATTATTTTCA TTACTGCATT TCATTGTAGG GAAGTAGATA ATTGCCCTTT
95281  ATTCACTGAC CTTCGCTTTT TAAAAATTTA AACCATGTTA CCATGAAAAT GCTTTTCAGT
95341  ATTTCTCTAC ACACAAGATT GCTGTAAGGG CAAAAATAGA GATAGGAATC ATGCATCCAT
95401  TGATATACAT ATTTTGATTT TTAATACATG TTACCAAGTT GCCTCCTGAA GGTCTGTTTA
95461  CACTCTCACC AACAGGGTGT TTTTTCCTGA CTTCCACAAA TGCTCTTGAA CAGTGGGTGT
95521  GTTAGTCTGT TCAAATTGCC GACATGAACA ATTAAATCTC ATTGTTGTTT TTATTTTTAA
95581  GACAATTATT GTTTGAGACT GCACATTTTG ATAATAACAT TTCTTCTATT ATGGTTTGAT
95641  TACTCATGAT TCTTGCCCAT TTTCTTTTGG GATGTTGCCT TATGTACATT ATTTTAAATA
95701  GATAGCTCCA TGTATTAAAA GATTATTAAG TTTGAGGGCT TATGATATGT CAGTTACATT
95761  TCTAAGATTT TTTTTTTTTT TTTTTGAGA CGGAGTTTCA CACTTGTTGC CCAGGCTGGA
95821  GTGCAATGGT GCGATCTCGG CTCACCGCAA CCTCCGCCTC CAGGGTTCAA GCAATTCTCC
95881  TGCCTCAGCC TCCCCAGTAA TTGGGACTAC TGGCAAGCGC CACCACGCCT GGCTAATTTT
95941  GTATTTTTAT TAGAGATGAG GTTTCTCCAT GTTGGTCAGA CTGGTCTCGA ACTGCCGACC
96001  TTGGCTTAAA AATCTACATT CTTTTTTTAA TTATAAAACT ACCACATCCC CCAAAAACAT
96061  TACTAAGGAA TTGAGGCTGC AGTTTAAGAA GCTGATATTT AGGATCTATC TCCGGAGAAG
96121  TGAGACCTGG TAATATAAGC ATTTTCAAAA TGAACTTTTG GGCCAGGTGA GGTGTGTCAT
96181  GCCTGTAATC CCAGCACTTT GGGAGACCTA GTCAGGCAGA TCACTTGAGC TCACAATTCG
96241  AGACCAGCCT GAGCAACATG GCGAAATCCA GTCTCTACAA AAAATTAGCA GGGCGTGGTG
96301  GCATATGCCT ATAGTTCCAG CTACTATAGA GGCTGAGGTG GGAGGATTAC TTGAGCCCGG
96361  GAGGCAGAGG TTGCAGCAAG CCAAGATCGC GCCGCCACAG CCTGAGCGAC AGAATGAGAT
96421  ATGCACCCAC GCCCTAAAAA AAAGCATGAC TCATTAAAAA AAAAAAATTT AGCCGGTCGC
96481  GGTGGCTCAC GCCTGTAATC CCAGCACTTT GGGAGGCCGA GGCGGGCGGA TCACGAGGTC
96541  AGGAGATGGA GACCATCCTG CTTAACACGA TGAAACCCCG TCTCTACTAA AAATACAAAA
96601  TAATTAGCTG GGCGTGATGG TGGGCGCCTG TAGTCCCAGC TACTCGGGAG GCTGAGGCAG
96661  GAGAATGGCG TGAACGCGGG AGGCGGAGCT TGCAGTGAGC CGAGATCGCG CCACGGCACT
96721  CCAGCCTGGG TGACAGAGCG AGACTCCGTC TCAAAAAAAA AAAAAAAAA AAAATTAAAA
96781  AAATATGAAG TTTTGAAGCA GAAATTATTT TGTCGTATGT TCTTTCATAA ATTTTTTGCC
96841  TGCCTGCCTT CTTCCTTTGT TACAGAACTC CAACACTTAC CCAAAGGTAG CTGTTGGGTC
96901  AGGGTTTCTG TACTATAGTC CCTTCTGTGG TGGCCAGAAA TATGTTACAG GAAAGAGGTC
96961  CCCATCCAGA CCCCAAGAGA GGGTTCTTGG ATCCCGCGCA AGAAAGAGTT CAGGGTGAGT
97021  CCGCAGTGCA AAGTAAATGC AAGTTACTA AGAAAGTAAA GTGGTGAAAC GACAACTACT
97081  CCATAGACAG AGCAGGACAT TCCCGAAAGT AAGAGGAGGA AGGCATCCAC CCTAGGTACA
97141  ATACTTGTAT ATATGGGGAG ATGTGCTCTG CTACAAGTTT GTGATAAAGG ATTAATTTTC
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 97201  TTAGTTACTA TATTTTGCAA GAATCAACAT TATTATCTTT AAACAAAATT AAGAATGCCT
 97261  TTGTTCTCCA GATATAGGGA TATCTGGACA CTCCTAAGTC TGAGTCTGTT TAGTAAACAT
 97321  TATTTATTTG TTCCCTTAAC CGTAAACATC TAGAAGCTAG GAATGACTGA CTTTCTGGGA
 97381  ATGCAGCCCA GAAAGTCTCA GCCTCATTTT CCTAGCCCTC ACTCAAAATG GAGTTACTCT
 97441  GGTTCAAGTA ACTCTGACAC TTTTCTTCTC TTTTTTTCTT CTTTTTTCCT TCCTTTATTT
 97501  TTTATTTTTT ATTTTTGAAA TAAGAAATCA AGAATACTTG ATGTTTCATC TAAAACAATA
 97561  CCCATAATTG ATAAGCCAAA ACAAAACCT AGGTCTTCTA ACTCAAAACT AGGATGTTTT
 97621  GCTGTCTCTG CTGATACTCG GCTGATCGTT AATAGGTAAT TAACAAACAA GCCTTGCTAT
 97681  GTCCCCCTCA GTTATTACC ATTAGATCAT ATGCCTACTG TCAATCATAT TAATCCACAA
 97741  CTATGCATTT CACAAAACTT GCCATAAAAA TTCACAGGTT TCCCGCTTCC CTCGAGTTTT
 97801  CATTTCCGAA GGGTCCCATG TAATATAAAA CTTATATTAA ATACATTTGT ATGCTTTTCT
 97861  CTTGCTAATC TTTTTTTTTG TTTTTTGAGA CTGAGCCTTG CTCTGTCACC CAGGCTGGAG
 97921  TGCAATGGCG CGATCTCGGC TCACTGCAAC CTCCGCTTCC CAGGTTCAAG CGATTCTACT
 97981  GCCTCGCCCT CCCGAGTAGC TGGGACCACA GATACGTGCC ACCATGCCCC GCTAATTTTT
 98041  GTATTTTTAG TAGAGACAGG GTTTCACCGT GTTGGCCAGG ATGTTCTCAA TCTCCTTACC
 98101  TCGTGATCCG CCCGCCTCGT CCTGCCAAAG TGCTCGGATT ACAGACGTGA GCCACTGCAC
 98161  CCGACCAATC TGTCTTTTTG TAGAGGGGCC TCAAGCATGA ACTTACTGAT GGGTGAGAAA
 98221  AACAGAATTT TCTTTTCCCC TACAATATAA ACATTAATTG TAATGTTATC ATTCAGGACA
 98281  TTTTGGTGAC CAATCTTACA GAAATTTTAT CTTGTCAAG TCTATGCAAA CCAATATGTA
 98341  AATCTTCTAT AAGTGAGATT GTATTCACT TTTCTAGTAT CCTTTTAAAT TAATAAAAGA
 98401  GATTCTAATG ATTATTTTCA TTACTGCATT TCATTGTAGG GAAGTAGATA ATTGCCCTTT
 98461  ATTCACTGAC CTTCGCTTTT TAAAAATTTA AACCATGTTA CCATGAAAAT GCTTTTCAGT
 98521  ATTTCTCTAC ACACAAGATT GCTGTAAGGG CAAAAATAGA GATAGGAATC ATGCATCCAT
 98581  TGATATACAT ATTTTGATTT TTAATACATG TTACCAAGTT GCCTCCTGAA GGTCTGTTTA
 98641  CACTCTCACC AACAGGGTGT TTTTTCCTGA CTTCCACAAA TGCTCTTGAA CAGTGGGTGT
 98701  GTTAGTCTGT TCAAATTGCC GACATGAACA ATTAAATCTC ATTGTTGTTT TTATTTTTAA
 98761  GACAATTATT GTTTGAGACT GCACATTTTG ATAATAACAT TTCTTCTATT ATGGTTTGAT
 98821  TACTCATGAT TCTTGCCCAT TTTCTTTTGG GATGTTGCCT TATGTACATT ATTTTAAATA
 98881  GATAGCTCCA TGTATTAAAA GATTATTAAG TTTGAGGGCT TATGATATGT CAGTTACATT
 98941  TCTAAGATTT TTTTTTTTTT TTTTTGAGA CGGAGTTTCA CACTTGTTGC CCAGGCTGGA
 99001  GTGCAATGGT GCGATCTCGG CTCACCGCAA CCTCCGCCTC CAGGGTTCAA GCAATTCTCC
 99061  TGCCTCAGCC TCCCCAGTAA TTGGGACTAC TGGCAAGCGC CACCACGCCT GGCTAATTTT
 99121  GTATTTTTAT TAGAGATGAG GTTTCTCCAT GTTGGTCAGA CTGGTCTCGA ACTGCCGACC
 99181  TCAGGTGATC CACCCGCCTC GGCCTCCCAA AGTGCTGGGA TTACAGGTAT GAGCCACTGG
 99241  GCCCGGCCAC ATTTCTAAAT TCTTTATAAG TATAAATTCA TTCAATCTTC ACCAAAACTC
 99301  AATGAAGTGT GAGTACTATT ATTATCATTG TTTTACAGAT CAAAACAAGT AATACAGTCA
 99361  CTTACTGAGT TCTATACACC TGGTAATTTT TTTGTTTCGT TGTTCTATCA ATTATTGGGG
 99421  AAGGGGTGTT GAAATCTCTA CCTTTAAATC ATGTATGTGT CTATTTCTCC TTTCGGTTCT
 99481  ATCAGGTTTT GCTACACATA TTTTGCAGTT CTGTTATTTG GTGCATATAC ATTTAGAATT
 99541  GCTTGTTTTT CGTATTGGAT TGACCCTGTT ATCATTATGT AATATCCCTG TCTGTTCCTA
 99601  GTAATTTTCT TTGCTCTGAA ATATACTTAT CTGATATATC ATCCAAAAGA CCACCAGGAT
 99661  GGCTAAAGAG TAGAAAGGAG AGATTTACTG GCAATACTAA TTTGCAAGCC AGGAAGAGAT
 99721  GGTCCCAGAA CCTGCCAAAA TTACTCTCTC TTTGGGGAGA AGGAGCAGGT TGGTTATTTT
 99781  TATGCCTCAT AGGCTATATA TTACACAATA GAGTCATACA TATTTAGCAC GTTTGGGGGG
 99841  ACAGCTATAT ATATTATGAG GGGTGCCAAG TGCATTCACA ATGGATAAAC ACGTGTAATA
 99901  TACCTCCCAT GTTCACTTCG AGGTTAAATT TTGGTTAAAA TGAGGTAGAA TTTAGGTCTT
 99961  TACATCACAA GGTGAACTAT AGGAACAAAG TTTACGTGCT GCCTCTAGCA GCTGGCTGAA
100021  AATGGCTTAA GGTCTACAAT TACGTGTAAG AATAGAATGT GTGTCAAGGC GGTCCTCTGT
100081  CCAATCAGAG TTGTAGTGGA CTGGACTGTA AATCAGAGTT AGGAGGGCTT CTGATAGCTC
100141  CTATAGTTAA GGAATTTAGC AAGTGTGAGT TTTTTGGTAG TCTTTGGAAT TTAGGAATTT
100201  GCCATGCCAG CCAAGCCATG AATGCTCTAC CAGTAGGTAA CTTTGTTTGC TTAATCTTAG
100261  AGTCTGTCTT AGTTGGTATA GGGGCATCTA TTTTGGTCTT TCAGATCCCA GATATTATTA
100321  ATACAGATAC TCTTGCAGTT TTGGGCTGAT GTTTATATGG CTTATCTTTT TTGCAGCCTT
100381  TAATTTCAAC CTGCGTTATG TTTATATTTG AAGTGAGATT CTTGCAGACA GTGTACAGTT
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100441    GTTGTTTTTT TTTTTTTTGA GATGGAATTT CACTCTTGTT GTCCAGGCTG GGGTGCAGTG
100501    GCACAGTCTC AGCTCACTGC AACCTCCGCC TCCTGGGTTC AAGGGATTCT CCTGCCTCAG
100561    CCTCTTGAGC AGCTGGGATT GCAGCCATGC GCCACCACAC CCGGCTAATT TTTGTATTTT
100621    TAGTAGAGAC AGGATTCACC ATGTTGCCCA GGCTGGTCTC GAACTCCTGA CCTCAAGTGA
100681    TCCGCCAGCC TCGGCCTACC AAAGTGCTGG GATTACAGGT GTGAGACCTC GCGCCCAGCC
100741    AAACTGTTTT TTTATGGGTG TATTTATACC ACACACATTT AATGCAATTA TTGATATCTT
100801    AGGGCTTAAG TTCATGAAGG GTAGTGTGGG AACCATAGTC TCTTGGCCCA CTAAATGTTT
100861    GCCAGAAATC ACTGACAAGG CAGATTGATT AATAGGTGAA AAGGCATTTT ACCTATTGTT
100921    TAACGTGTCT ATGTGGGAGC ATTCAGAATT AATTACCTAA CTTCCCAATG AGTTATAGAT
100981    GCTTATATAC CATTTTTAGA TCACAGAAAG AATTGGGGCT TAGATTCTGG TAAAACAGGT
101041    TATGGGAGGC AAAAGAGGTT TGGCTTGCAA AGGTGGCCTT GTTAGGTAGG TGAAGCCTCC
101101    CTCAGAAAGA ACAGATGGTA AATGTTTCTT TTATGATTTT TAAGTGTCAG ACTCTCAGTC
101161    TCTCCTGGAT CTGGGGAAAG GTATAGAAAG GTGAGGAGGC ATGGCTGCAT TAATGGAGAT
101221    TCTCTACAGA TGTAAAATTT TTCCCATTTA AGGCAGCTTT GCAAGCCCAT TTCTGCCTGC
101281    TGGCCAAGCA GCAGCCATTT CAAAATATGT CAAAGAAATA TATTTTGGGG TAAAATATTT
101341    TGATTTCCTT TAGACTGGTG GCCTTATAAG AAAAGGAAGA GACACCTGAG CTGACACACA
101401    TACCCTTGCT CTCTCAACAT GTTATGATGC AGTAAGAAGG CCCTCACCAG ATACTAATTC
101461    CATGCCCTTA GCTTCCCAGG TTCTAGAACA GTAGGAAATA AATTTCTTTT CTTTAAAAGT
101521    TAGCCAGTCT GTGGTATTCT GTTATAGTAT CACAAAATGG ACTAAGTAAC TATATTATGA
101581    TCATCTTACA TGACTGATCC CTCCTACATC ATACACATAC ACAGGCCACA TTTGGAACAT
101641    TGTTAGAGGT TCCTCTACCC AGTACAAATG TACTACAAAT TATATATGTA TTTTTAAATT
101701    TTTGAGTATC TTCAATAGTA TATTTTCGTT AACTTTTGTA GTCAAAATGT CATTATAACA
101761    TGTATTCAAT ATGCATAATT ATTAGTCAGA TGTTTTACAT TCTTTCTTCA TACTAAGTGA
101821    TATGGTTTGG ATATTTGTCC CCTCTAAATC TCATGTTGAA ATGTAATCTC CAATGTTGGA
101881    AGTGAAGCCT GGTGAAAGGT TTTTGGATCG TGAGGGTGAA CCCCTCATGA AGCGCACTCT
101941    TCAGGGTAAT CAATGGGTTC TCACTTTGAG TTCACAAGAG ATCTGGTTCT TTAAAAGAGT
102001    GTGACACCTC CCCCATCTCT CTCGCTCAGC TCTCACCATA TGATATGCCT ACTCCCTCTT
102061    CACCTTCCAC CATGATTGGA AGTTCCTGA GGACTTGCCA GTAGCAGATG CCTGCACCAC
102121    ACCTCCTGTA CAGCCTGCAC AACCGTGAGC CAAAAAAAAT TACTTTTCTT TATAAATTAG
102181    TCAGTTTCAG GGATTCCCTT ATAGTAATGC AAGAACGAAC TAACACACTA AGTCTATTTC
102241    ATATTTACAG AATAGCTCAA TCTGAAGTAC CCTTTTTCAA CTTCACAGTA GCTACTTGTA
102301    GCTAGTGGGC ACTGATTTGG AGCGTGTTCA AGGGTGAATT GTATTATGCA ATTAACAGAT
102361    TTTTTTTATT GTTTTCGCAA ACCACGAGGC ATAGATTGTC TTACTTTCTC TGCTCCTGGT
102421    GTTGGAGTTG TTATTGGGAA ACAACTTATT TTCCTCTTAT ATTTATATGG AATAAATAAC
102481    CCCCAATATT TCCCTCCCCA ATATCTGCCT TTTGTATGTT TTTTGAAGGC AAGTGCCTAG
102541    AATTTACTGT TTTTGAAGCA CTTACTGAAA GGATTGCCAT CAAGTTGTTT TGCTAATAGT
102601    ACATGCCAGG CGCTTGTTGG TTTGCTTAAT TCAAGGTAAC TTGGATGAGA AGAAGAGTTT
102661    TTCTCATCCA TGGCTCAGTG GAGTATAGAT TACTGATATT GTGACTGGAT GTACTCCTGC
102721    TTTCTAGTCT GAGTTTTTGA AGCTACCCTT AATCTTGGTT TCAATTTTAT CTAGCCCTGT
102781    ACATATCCAA GGCTCTTTCC AAAATGGTCT ACGATTTGTT TAGGAAGTTA GAATAGCTGT
102841    ACTTTCTGAA CCACGGTTCC TGACATTTTC TGGACTTCAA ACACATCCAG CATTTTATCG
102901    AAGTATTTAT CCTTCCTACT TGGCTGGCTT CTTCCTTGCC TTCAGGTCTG AATTCAAATG
102961    ACATTCTCCT GATGAAACTT TCCATCCTTA TTTCTATTCT TTTTTCTTAT CCCCTTTCTT
103021    TATTTTTCTC CACAGCACTC ATCACTTATC TCTACATTTT CATTATGTAT TTACCTTATT
103081    GTGCACCTCC CACTACAAGA CAAGTAGCAC CGTAAGGAAA CAGGTTGTCT GCTTTTTCAC
103141    TGCTATGCTC CCTGCACCTA GAACACTCTC TGGCACTTAG CAGGTTTTCA GTAAATATAT
103201    GCTGAACTAA TAATGCTGGA TATACATCTC CCTCATGAAC TCTCTAAATC CTTCTAATTT
103261    ACATTGATCA ATCTTCTTTT CCATGTGCTT TTGTATGATT TATTGCTCAA AATCTTTATT
103321    TTGTATGCAG AACGTGCACT GCTATTTAAT CTTCATGTAC GTAAGTCCTC CCTTCTCTGA
103381    GTATAATCTC TTCAGGGCAC TATCTGAGAT AACTTTTTAA CATCTCCATC ATGAATCTTG
103441    TACCTTTTCA AAGAAAATGA GCCAGTGATT ACTGATGTTT ACGGCTATTG TTGAGGGTGA
103501    AGATCATTAT AATTTTGAAA AGGGAAGTTG AATATTGTGA AGGGAAAGAT AACACTAGAG
103561    TCAGAAGACT TGGGAGAAGG CAAAAAACAA ACTAAAAATG AGCACTTTTA GTCTCCTGAC
103621    AGTTTCTCTG AATCAAATCC ATAGTTCTGT GACAGCGTTG GCTTAGAAGC AGATTTTTTT
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103681   TTTTTTTTTT   TTGAAATGGA   GTTTCGCTCT   TGCCCAGGCT   GGAGTGCAGT   GGCACGATCT
103741   CGGCTCACTG   CAACCTCTGT   CTCCAGGGTT   CAAGCGATTC   TCCTGCTTCA   GCCTATGGAG
103801   TAGCTGGGAT   TACAGGCTCC   CACAACCACG   CCCAGCTAAT   TTTTTGTATT   TTTAGTGAAG
103861   ACTGGGGTTT   CACCATGTTG   GCCAGGCTGG   TTACGAACTC   CTGTTCTCAA   GTGATCTGCC
103921   CGCCTTGGCC   TCCCAAAGTG   TTGGGATTAC   AGGCATCAGC   CACCGTGCCC   AGCCAGGAGC
103981   AGATTTTTTT   ACACTCATGT   TTCTTTTTCC   TTCTGTCATC   CTGTTTCAGT   ATAAGCAGAC
104041   CACAGATAGA   AGTAGTAGAT   ACCTCAGAAA   TTCCTGGAAT   AATTAATCCA   CGTTCATCTG
104101   TACTCCATCT   GCTCCTATCT   CATGGAATAT   AAAAGGAAAA   ACACCAAGAT   TTCCCTAGGC
104161   AATCTGTCTT   GATTTTAGGT   TCCTCAACAG   GAGAGCCAGA   CAATGGCTGT   AATAATATTG
104221   TCCCGGCCAA   GGAAAAACTT   CCCCTTTGCC   CTCCCAAGGT   TTATGGAAAA   TTACTGGCAA
104281   AACACAGATT   AACTGGAGAA   AAGGCATATA   TATTTATTTC   ATCACAATTT   TACAGGAGAT
104341   TTTAGAATTA   AGACTGAAAG   ATACAGGGGA   AATTGCCCAT   TTTTATGCTT   AGGTTCAACA
104401   AGATAAACAG   CTGTATAGGG   TACGATCTAA   TGCTAACAGA   CTGAGTGGGG   AAGCCCCGCA
104461   AGGCTTGTCT   GTCAAGATTC   TTCTTGACCT   CTCAGTGCAG   CATTTCTTCC   TTCTGGTTAT
104521   AGGACAAGAC   TCTCTTTTAG   AATGGGGGGT   CTTATGACCT   ACAGGCAAAC   AAGGTAGGTT
104581   AGAGTAATAT   TTTTAGGTTT   TATGGCTGGT   TCTAGGGAAA   AGGAGTTCTG   GTTTGTATGG
104641   CCTACCTTGA   GGAGGAATTC   TGGTTTCTAT   GGCTAGACTT   TGGGGAGAAT   GGGACTTACA
104701   GACAGGAAGG   CAGAAGGTGG   TCAGTGAAAC   ACTTTTATAA   TCATAATCCC   ATTTTGAGTA
104761   TTTCTGTGTT   ATGGAATGTT   TGTTCTCTCA   TTTCCTGAAA   GATTCCAGAG   ACTCCTCATT
104821   CAGTGTTGTG   AAAAAGTTCA   GGAAATGCAA   CTCAAAAATG   TGCCACTTTG   TTACGCTGAT
104881   TTCTTTGAAC   TGAGGGCACC   TAGGAAACAG   TAAATTCAAG   GAAGGGCTTT   CGCTGAACTC
104941   TAATCAAAAA   TTTGAAAATT   AAAAAAAAAT   TCAAAAGGA   ATTTAGTTGT   TAAGATTCAC
105001   TTCCCTGGGG   AATCTCATCA   ACCAGAGAAG   ATTAACTGTA   TCACAGGAGA   GGAGACTGGT
105061   GGTTAACACC   ATCTAAACAG   ACTTTGTCAC   AGCTGTCACC   TATTCTTTGA   AACACCCATT
105121   TATTTTTCTC   CAAAATCATA   TACTCTCCCC   TAAGTTGCCT   ACATCCCCCT   TCTTTCTCCC
105181   TTATGAATCA   AGAGAGCTTA   TAAGCTTCTA   CAGTTCACTG   GGATTTGGGG   TATTCGCTTT
105241   TCTTCCCTCC   CACTCCCCCT   CCCCTTTTTT   TGTCTTTGAG   ACACAGTCTT   CTGGCTCTGT
105301   CGCCCACGCT   GGAGTGTGGT   GGCTCTATGT   GAACTCACTG   CAACCTCCTC   CTCTCGGGTT
105361   CAAGCGATCC   TCCCACCTCA   GCTTCTCGAG   TAACTGGAAC   TACAGGCGTG   CACTACCAAG
105421   CCCGGCTTTT   TTTTTTCTT   TTTCTCCCCC   GTTTCTTTTT   TGGTTATTTT   ACTGGAGACA
105481   GGGTTTCTCC   ATGTTGTCCA   CGCTGGTCTC   GAACGCCTGA   CCCGCCGTCC   TCGGCCTCCC
105541   AAAGTGCTGG   TATTACGGGC   ATGAGCCACT   GCGCCCGATT   TGAAGGACCT   CTTAAATATC
105601   TATTTAGAAA   TTGGTCGGAG   TCCACTCCTT   TCCAAAAACA   TGAGTCACAA   TCCGGGAAAA
105661   GCACGAGCGG   CTGAAAGTCA   AAATAACCAG   AACAAAACCT   CCACTCATGC   TTAAAAAAGG
105721   TATTTTGACA   AAATCCTAAT   TCGGCCAATT   ATTATTAGTA   TTCAAGTCGA   AGGCTCGTCA
105781   AGCCAGACTG   GGGATTGGGT   CAAACATAAA   CCTTACACCA   GACGGAAGGA   TTACATGCAA
105841   ATGAAGGATG   CAGATTCTGA   TTTCCCATTG   GGTATTTGAC   ATTAGCCAAT   GGGAGAATTC
105901   CTCACAGCCT   ACCTCCAGTC   AGTATAAATA   CTTCTCTGCC   TTGCGTTCTA   ATGTAGTTTC
105961   ATTACATTTT   CTTGTGGCGA   TTTTCCCTTC   TTATCAGAAG   TAGTTATGTC   TGGTCGCGGC
106021   AAACAAGGCG   GTAAAGCTCG   CGCCAAGGCT   AAGACTCGGT   CTTCTCGTGC   AGGTTTGCAG
106081   TTTCCTGTGG   GCCGAGTGCA   CCGCCTGCTC   CGCAAAGGCA   ACTACTCCGA   GCGCGTCGGG
106141   GCTGGCGCGC   CGGTGTATCT   CGCGGCGGTG   CTTGAGTACC   TGACCGCCGA   GATCCTGGAG
106201   CTGGCGGGCA   ATGCGGCCCG   CGACAACAAG   AAGACCCGCA   TCATCCCGCG   CCACCTGCAA
106261   TTGGCCATCC   GCAATGACGA   GGAGCTTAAT   AAACTTTTGG   GGCGTGTGAC   CATCGCGCAG
106321   GGTGGCGTTT   TGCCTAATAT   TCAGGCGGTG   CTGCTGCCTA   AGAAAACTGA   GAGCCATCAT
106381   AAGGCCAAGG   GAAAGTGAAG   AGTTAACGCT   TCATGCACTG   CTGTTTTTCT   GTCAGCAGAC
106441   AAAATCAGCC   TAACAGCAAA   GGCTCTTTTC   AGAGCCACCT   ACGACTTCCA   TTAAATGAGC
106501   TGTTGTGCTT   TGGATTATGC   CGCCCATAAA   GATGTTTTTG   AGGTGTTTTT   AATGGCTTTG
106561   AGTGTGGCAC   TTTTAGTAAT   TTGTCCTGCA   GAAATTAGAT   CCATAGAAAC   CTCAGGAATT
106621   CTAGGTATGT   GGGAGAAGTG   CCATGCAGCA   CAAAACATGT   TTACAGGGGT   GATTCGCGTT
106681   AAGTTTCACA   CACAGCAGTT   ACTACATTTT   AGAGGAAGGA   AATTATACCC   ATGAGTGCAT
106741   TCCTAACTAT   CTTGAATGGA   AGTGTTAAAA   CCCGCATGCC   CCACACAAGT   TTGAATATGT
106801   CATACCATTT   GCTGTAGCAA   TTAATGGCAT   ACACAATTGA   GAGCACACAC   ATTACCACTG
106861   AACATTTGAG   TATGTATTTC   CCAAAATGAG   CTTTTTTCCA   GTTTGGGGAT   GTTTTGCTTT
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106921   GTTTTGGGGT GGAGTCTCCC TCTCGCCCAA GCTGGAGTGC AGCGGCGTGA TAACAGCTCA
106981   CTGTAACCTC GAACTCGGGC TCAAGCGATC CTCTTGACAG CCTTCTGAGT AGCTGGGATT
107041   ACAGGCGAGA GCCGCCACGC CCGGCTAAGA GCATTTTTCT AATTGCCCAC ACTTCTTATG
107101   CGACACCCAG AAAAATACAA TTTTAAATAA AGCGCATATG CAAATTTCCC TAATCGTCTC
107161   CAATATTCTC TGATTTCTTT TTTATATTTT AACTAGAAAC AATTGGAGGT TTCCGCGTTG
107221   CTTTGTGTGG TTGTAAATTT TAAGACTTCA GGAAACTTTT CCAGTACAAG ACTTGTCCAC
107281   AGTGGATATA GCAGCTAAGG GGTTAACAAA ATGACGTCAG AGTAGCTACG GTAATGGGCA
107341   GGAGCCTCTC TTAATCTGCA ACCAGGCACA GAGATGGACC AATCCAAGAA GGGCGCGGGG
107401   ATTTTTGAAT TTTCTTGGGT CCAATAGTTG GTGGTCTGAC TCTATAAAAG AAGAGTAGCT
107461   CTTTCCTTTC CTCCACAGAC GTCTCTGCAG GCAAGCTTTT CTGTGGTTTT GCCATGGCTC
107521   GTACTAAACA GACAGCTCGG AAATCCACCG GCGGTAAAGC GCCACGCAAG CAGCTGGCTA
107581   CCAAGGCTGC TCGCAAGAGC GCGCCGGCTA CCGGCGGCGT GAAAAAGCCT CACCGTTACC
107641   GCCCGGGCAC TGTGGCTCTG CGCGAGATCC GCCGCTACCA AAAGTCGACC GAGTTGCTGA
107701   TTCGGAAGCT GCCGTTCCAG CGCCTGGTGC GAGAAATCGC CCAAGACTTC AAGACCGATC
107761   TTCGCTTCCA GAGCTCTGCG GTGATGGCGC TGCAGGAGGC TTGTGAGGCC TACTTGGTAG
107821   GGCTCTTTGA GGACACAAAC CTTTGCGCCA TCCATGCTAA GCGAGTGACT ATTATGCCCA
107881   AAGACATCCA GCTCGCTCGC CGCATTCGCG GAGAAAGAGC GTAAATGTAA AGTTACTTTT
107941   TCATCAGTCT TAAAACCCAA AGGCTCTTTT CAGAGCCACC CACTTATTCC AACGAAAGTA
108001   GCTGTGATAA TTTTTTGTTG TCTTAACAGA ACAAATTTCT AAGGACCCCC CCGGAAAGCA
108061   TTAGACTATG GTCTTAAAGT TGATTAACAG AAATAACGGT TTGGTCAGTC TTGCAGTGTA
108121   GGTTATTTCT GACCTTATTA AGGTGCTATT TGGAGAGAAG CTGTGTAAGT CCACTATCAT
108181   TCAGGCCTCT AGCTTGCTAT GATTAGCATT TGTTTAAACA ACTTTGTAAG AGTAAGGGAA
108241   AAATCTGGTA AGTAGTTAAC TGGCGCTTAC TAGGCATTTT TGCAAAGCTT TGAAAAGATT
108301   AGAAAATTGT GTCTTGCGAG TTCCAGTGTC TTCCTCAAAA TGCTTAGGAA GATTTTCTCA
108361   GCTCAATACA TAGTCCCCTA GGTTTTCTCA TATATTATAT ATATATATAT ATATATATAT
108421   ATATATATAT ATATACTGTT AAATTCATTT GGCTGTTAAC ATTAACCTGA AATTTATTCT
108481   GGTGCAAAAT GTGAGGCAGG GATCTAACTG GCTCTCATTT TATCCATAGC TAGCTACCCA
108541   CTTTAAATCT GTCAGTCTGT CGACCAAGCA TAATTTAATC CCTTATATAT GAATTTTTAT
108601   ATGTGTGGCT TTGCTTGTAA ATAGTCTATC TGGTTGCATT GCTTTGTCTC CTCTAGGACT
108661   ATGCACCATG ACATGCCACA TTCTTTTTTT CAGTACTTCT TGCCTGTAGT TATTAAAATC
108721   TAGAATTTAC AAGTTTTAAC CATTTTCTTT CTGTTGATCT TGCTTTTCGG TTTTGGAGGT
108781   TGGGGATTGA GTACTGGAAG AAAATTTAGA GGGATGGGAA TACTGTACGC AAACAAAAGT
108841   AATATTTACT TTAAAATTTT TATATTTTGT ATTTTTTTAT CATATAGCTT TTACATCACA
108901   TTTTACAGAC TAACTTTAGA ACAACCACAG AATGTCCAAC ATTAAAACTA CTAATTCCAA
108961   AGACCTTGCC TCACATTCTT TTTTACAATA AATATTTTTT ACACCTAACA TTCTTTCTTG
109021   GCCTACATCT AGAATGTAAA CTGATGTACC ATACTAAAAT CGCCTGACCA ACTGTCAACA
109081   ACAACAAATC ACACACACAA AAGATCAAAT TTGAATTGCA TCGTTTACTT AAATTCATTT
109141   GTGTTCCAGC TTTTAATAAG GCAGTTTTTG GTTTATAAAG TAATATTTGC ATTTTAAAAA
109201   TTATGAAAAT GAATATGTCA GTTTGTTTTA TGATTCGTTT TTCTTGACTC TTATACAAGC
109261   GACTCTAACT GGCATAGACA TTTGTTATCC ACAGACAGTA TAGATATGTT AGAGATGCCA
109321   ATGGACTTGG TCTATGCCAA GGTGACTACT CACAAGCTCT GGGCCCAGCT GAAGGTCAAG
109381   TATTTTTTTT CCAGTTATAG ATGTGCTGGA TCTGATGTAT AGCGCTTGAC TTTTTATATT
109441   TTCTTTATCT GTAGGAAACA AATGTGTTGG AGGTACTGGG TCTGACGAAT AGCATAAAAG
109501   AATAAAGTTA CATTACTGTC TGAGGATCAG ATGGACAGGG GGTGGTAGCT CAGTCCAGCT
109561   ATTTTCCACT CCCTCACTTA CATTCTTTGC CCCCTCCTCA ACAGAACAAG GATTCTGCTG
109621   TAACTCTTCA TTGACAGTTG ATATTTAAAA ATTAACGAAT GGATGAAATT CTCATTTGTG
109681   AAAGAAAATT TATTGAGCAT TTTGTATTTG TGAGTAGTGC AAACATTTTA ATATTATATT
109741   AAGAATCTAT TGTTTTGTAT TAGAGGAGTA ATTAAGGAGA GATTGGAGAC AAAAAGGGGG
109801   TGTTGTTTGC AGAATATACC ATCCAAAAAT AGACCACTGT GGGATCAGGA TTCTTTTGAG
109861   CTAAAGGCAC TTCAAAAACA GCATTCAAGA AGGGAATTCT TCTAAACTTT TCTTTCTGAA
109921   AACAGGAGAT AAAAGTTCCA ATGTGAAAAA TGCTCTGCTT GTACCAGGTG AAAAGACATA
109981   TTCTTCAGCC CAGAGGCATA GATGAGATAA TTCTGCACAA ACACAGCAGG GAGTCATAGC
110041   CGAGAGACTT CTATACACAA ACAAACCTTG TTAAAATAAT CATATATTCC TTTAATCTCC
110101   TCATATGGTT TACTTTCCCA CAATTGCCTC TCTTTAACTT AATGTGAAAG CATTTAGCTT
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110161  TTGCCATTTC  TTTGGGGCTT  CACTTTTTTA  TGAGGGTTCT  CCTGTCCCAT  AAAATTTACA
110221  TTAAATACAT  TTGTATGCTT  TCATTCTGCT  AATCTGTTTT  ATGGCAAATG  AATTATCAGG
110281  TCCAGCTGGA  GACCCTAACA  GAGTAGAGGT  AAAATTTTGC  CTCCCTACAA  GATAGAGATT
110341  GTGTGCATTA  AATGTTGTTT  GTTCCCAGTT  GTTCAGTTTG  TCAGGCCTCT  GAGCCGAAGC
110401  TAAGCCATCA  TATCCCCTGT  GAACTGCACG  TATGCCTCTA  GATGGCCTGA  AGTAACTGAA
110461  GAAACACAAA  AGAAGTGAAA  ATGCCCTGTT  CCTGCCTTAA  CTGATGACAT  TACCTTGTGA
110521  AATTCCTTCT  CCTGGCTCAT  CCTGACTCAA  AAGCTCCCCC  ACTGAGCACC  TTGTGACCCC
110581  CACCCCTGCC  AGCCAGAGAA  CAACCCCCTT  TGACTGTAAT  TTTCCACTAT  CTACCCAAAT
110641  CTTATAAAAC  GGACCCACCC  CATCTCCCTT  CGCTGACTCT  TTTCGGACTC  AGCCCGCCTG
110701  CACCCAGGTA  GAATAAACAG  CCTTGTTGCT  CACACAAACC  CTGTTTGATG  GTCTCTTCAC
110761  ACGGACGCGC  CTGAAACAGT  TTAACAGGGT  TTTTCCTGCC  CAGTCACAAC  AAAGTGATGT
110821  TATGCTGCAG  GCTGAAGTTT  ACAGCTAATG  CTGTTGAAGT  CTAAAATCAG  TTTTGGTTTG
110881  TTAGATTTGG  GTGAGATGGC  TAAGATTCTC  AGAGAAAGAA  GTCAAGTTTG  GGGTGCATTT
110941  TTCAGACTTA  AAAATTTAGC  AGTAGCCCTT  GCAGTTTTTC  CAATAGAAGT  GATTTACGAA
111001  TGTTTTCAGG  AAATTTAAAA  CAACAGTGAG  AAGCGTGTAT  GGAGAGTTGA  ACTACACTCC
111061  AGACTTGGCT  ATAGGAAAGC  ACGAATGCTG  CTATTGTATT  GCACCTTGGA  AAAGAGAACA
111121  AAGGAATATT  TTCGGACAAT  TTTAACATGT  CACATATGAA  AAGCTAAACG  GAATCTGTCA
111181  ACACCTTGTA  CGTTATTACA  GGCTGTGATT  TTAAAAAAAC  AATCCTTACT  AATACATACA
111241  TAGTTGCTGC  TAGCAATATA  GTGTTGGGAG  TAAAAACACG  AAAATGAGAG  TTCAGGACAA
111301  TATCCCAACT  CTGAGCAGAT  TTTTTTAAGT  AGTAACATCT  AAAATTAAAC  CATATTATGT
111361  AATATTTATT  TCTTTTCCAC  AGTCTCTTCT  CATGCCTCGT  TCACATTAGC  TAATTAAAAG
111421  TCCCCTGAGT  ATCATCATAA  CCCGATTTAC  AGATGAAGGC  ACGGTTGCAA  TGAGCTATCA
111481  CCCTCTTCTG  AATGAGACAG  TACAGTGTGA  AGGATAGCAA  AACTCCACTC  CCATCCTCTT
111541  AGGGCTCTGG  CTGGACCAGC  AAATTAAATT  AATGTAAAAT  GGATTAACAG  GAGAAAGGTA
111601  TATGCATTTA  TTTAACACAG  GTTTTACGTG  ACACAGGTGC  TCTCATAAGG  TAATGAAAGC
111661  CCAAAAAAAG  CAGTTAGCTA  CTTATATAAT  GAATTGGACA  ATTAGTAAAA  TGTAAAAATG
111721  CGCTAAAGCA  AAGGGATTTA  GGCTAGAATA  TATAACTGTG  TAGAGAAGCG  CCCAGCAAGG
111781  GCTAGTGCAA  GGTTTGTACA  GAATTCTCTT  GGCCTCAGCC  TCCTATCCTT  GAGAAGAATG
111841  TTGCTTTTTT  TAAACTACAG  TGAGAACATC  TTTCATATGA  GAATTTCACC  TACTGCTTCT
111901  AAGAAACAGG  TCAGCTTTCA  AGAAAACATA  AGGCCAGAGT  GATCTTTTCA  CGCCTGCTCT
111961  TTTAAGTACC  TTTGAATAGT  CAATATGTCT  TCAAGCACTT  GAAAGACTTA  AAAAGTTTAC
112021  CACTCCGGCA  TATTAGTGAA  AGCCCTTAAT  ATAAGCCCTT  ATTAAAATTC  TCAGTCGAGG
112081  GTATAAATTC  AGATTCAAAT  AGTAGTGTCG  TAAACGGGAG  GGAAAAACTA  AAGGGATTAA
112141  AAAGTGAAAC  TATTGTGTTC  TCCCTCGCAG  TCCTTAGGTC  ACTGCCCCTC  GAGGGGCGGA
112201  GCAAAAAGTG  AGGCAGCAAC  GCCTCCTTAT  CCTCGCTCCC  GCTTTCAGTT  CTCAATAAGG
112261  TCCGATGTTC  GTGTATAAAT  GCTCGTGGCT  TGCTTTCTTT  TCGCGTACCT  GGTTTTTGTT
112321  GTCAGCTGGT  TAGACATGTC  TGGTCGCGGC  AAAGGCGGTA  AAGGTTTGGG  TAAGGGAGGT
112381  GCTAAGCGTC  ACCGAAAAGT  GCTGCGGGAT  AACATCCAAG  GCATCACCAA  ACCGGCCATT
112441  CGGCGCCTTG  CTAGGCGTGG  TGGGGTTAAG  CGAATTTCCG  GTTTGATTTA  TGAGGAGACT
112501  CGTGGCGTTC  TCAAGGTGTT  TCTGGAGAAC  GTGATCCGGG  ACGCCGTGAC  CTACACGGAG
112561  CACGCCAAGC  GCAAGACTGT  CACTGCCATG  GATGTGGTTT  ACGCGCTCAA  GCGTCAAGGA
112621  CGCACTCTGT  ACGGCTTCGG  CGGTTAATCT  TTTCGTCAGT  TTTCTTCCAA  TGGCCCTTTT
112681  TAGGGCCGCC  CACTCCCTCT  CAGAAAGAGC  TGTGATTGTA  TTCTTTCGGA  TGGTAACATC
112741  TCAATGGCTT  TACTCGGCTA  TTCTGCCTAG  TATGTAGAAC  TATTATAAAC  CAGTTGGGAG
112801  AGACCAGGTT  GTTTGGTCTG  AGTGGCTGCT  AAAGCAGAAA  TCAGCTAAGT  AAACGAGGTC
112861  TCCGAGATAA  GTGAGCTATA  AACTTCAATG  CTATAGTTTT  GACATGTCAA  GCAACTTAAC
112921  GTGCAGCGCG  AGTCCGATAA  ATGAGTAGCT  CAGCTTTTTA  GTTTTAAAAA  CGAGTTGTGC
112981  GTTATTTGTA  CGAGAGCCTA  AGATGCTAGC  TGCCTGGAAC  TGAGTAGGTG  GATTAAAATG
113041  GGTGTCAGGT  CTGTTTTCCC  AGGCGTATCT  GACTTAACGT  CAGCAAAAGC  TGTACTTTTA
113101  GCTTCCCTGG  TAACACCTGC  CGTCCTTAAC  CGCCCCCTGC  CGGTAGCGCC  AGAAGCCTTT
113161  ACTTCCATTT  CTAGTTGAGC  TTGGCGTCCT  GCTGAGTGAC  GTCACCTCCC  CCTTCTGTGG
113221  AGTAGGACTG  GCGGTTAAAG  CTGCTTTGCT  ATTTTCAGTC  CTCAGGCTGG  AGGCTCCCCT
113281  AAGCAGGCTG  CCTACGCAGT  TCGTAAATTC  CCACTTAGTA  GACTAAGGGA  GTCTGTTTTA
113341  TAAATAAGGA  CTCAAATTTC  TTCTGACTCC  GAGGTCCGTG  GCAGCAGCTA  TAAGATGGAA
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113401  GCCCCCTCTG ATGTAAGATT CTCAGATGAC TTGCATCTTC ACTGTACCTG TCAACCCAAT
113461  AGTCTTCTAT TCCTGCCTTA AATTGTAAAT TCCAAAACTG ATTTAATTGT GAAAGTTTCA
113521  AACTGTACGA CCTAGGAAGT GTCAAAGTTA GGTGACCAGA TTTTTAGAAG TCAGCCAAAT
113581  ATTCAGCATC TTTGATTTAG TAACAAATAT ATTGATGGCT ACTTCAGCAA AAAAAATCAA
113641  CTTTGTTTTC TGGTTACTTT GCTAACAAGC TTCTCCTGAC AGGAGGATAT AGTGAATAGG
113701  CAGTTGAATA AGTGAGTTCG GGTGAGAGGT CTGAGCTGGA GATAAAAATG TGTGAGTCAT
113761  CAGCAGATAA ATAAATGCTG AGACCAGATG AGATGGCTAA AAACTGAAAC ATAATGTAGT
113821  GCAGCATTGT TTGTAATAGT AAATGAGTGG CAACTGTAAA GTTTTCATCA GAAAGGACTA
113881  GAGTGATCTA TACATCCATA AAATAGAGTA TTTCTCTACA CAGCCCTACT AAAGAATGAG
113941  AAAGCTGTAC TCCACTACAT ACTCTGGTGT ACTCTGGCTC AGTTCTTGGA CTCCTCTTTT
114001  CTTGGCTAAC TCAACTGGCC TCACCACTTA CATGCTCTGT GCTCTGTCAA ATAGTTTGTT
114061  CAACAGAACA CCACGGCCTA GCTGTAAGTG CCACGTTAAC TTCTAGCAAT GCCAAAGCCT
114121  GTGATAGTGG CAGCTTCGGG CTGTTTCTCA TTCCCGGGAT GCCTAACCAC CTCTCCAAAT
114181  TCTATCAGTT TGCTTCCACC CACTTCAAGC TTCAGAACGA AACATAGAGC TTAAGAAATA
114241  TAGGCCCGGC AAGGTGGCTC ACGCCTGTAA TCCCGGCACT TTGGAAAGCT GAGCCTGGTG
114301  GATCACCTGG GGTCAGGGGT TCGAGACCAG CCTGGCCAAT ATTGTGAAAC CCCGTCTCTA
114361  CTAAAAAAAA AAAAAATTA GCTGGGCATG GTTGCGGGCG ACTGTAATCC AAGCTACTCG
114421  GGAGGGTGAG ACAGGAGAAT AGCTTGAACT CGGGAGGCAG AAGTTGCAGT GAGTTGAGAT
114481  CGCGCTATTA CACTTAGGCC TGGGAGACAA GAGTGAAACT GTGTCTCTAA ATAAGTGTTT
114541  GCAATTATAA ACCATCTCCC TGACCTTAAA TCTCTAGACT CATATACAAC TGCATATTTG
114601  ATGTATCTAA TTGAATAATG GGCATCTCGA ACTTGTCCAA AATATGTTTA TACGTAAACA
114661  CCAAGTCTGT TCTTCCTCTG ATATTTGTCA TGTCAATCAA TAGAACTCCA TTCTTCAAGC
114721  AGCTTGGGCC AGGAATTGTG CAATATTGTT TGTCCTGAGC TTCTTACAAC TTTCACCCAA
114781  TGCAGTCAGC TCTGTTGAAA ATCAATCAGA ATACCTTTCA TTGTTTTCTT TGCTGCTTCT
114841  CTAGGAGCAA GCTGCCATGG CGGTTTGTCT GAATGACCAC AGTGACCCCA AACTGGTCTT
114901  TGTTTTCACT TTTAATCCCC CTGTCATACA GTTTTTCTCT ATCCAGCATC AACAGTGATC
114961  CTTTTTGAAG GTATTATGTC CACTGTCTGC TGAAAAGATT CCACTGGCTT TCCATCACCT
115021  TCATAATAAA AACCAGCATC CTTATCATAG CCTACAAGTA AGATGACCAA CCATTACAGT
115081  TTGCCTGACT CTCAGGGGTT TCTCAGGGTG TAAGACTTAC AGTGCTGAAA CTTAGAAAGT
115141  TCCAAGCAAA CTAGGATGAG CTGCTCAACC TACTAGATCT GTACTCTGGC TACCCTCTGA
115201  CCTCATTCTC TTCGCAGTTC TTTCTCTTCA CTGACCTTGC TGTTTCTGGA ATGGACCAAG
115261  CATTTCCAGC ATCAGCACCT TTATATCTAT TCTTTCTCCC TAGAAGGGTC TTGTCCTGGA
115321  TATCTGAATG GCTCTAGATC TCATTTCATT CAAGCCTCTC CTCAAATACC AACCTTAAGA
115381  AAGAGACCTC CCATAATCAT CCCTTGTAAA ATAAGCTTTT CTGCTCATTT AGCATATATA
115441  TATATAGTTG ACTATCCTCA ATAGCATATA TATATAACAT TTCCCCACCT AGAATTATAT
115501  ATGTAATAAT ATATTTAACA AAAAATACAT ATAACTAGAT ATATTTTATT TTGTGTTTGT
115561  TCTCTCTCCC CCAACTGGAA TATATTTTTT GAAGGTAGGG ACTTTGTTTT GTCCCAGAAG
115621  TATCCCTAGC ACCTTGAACA GGGCTGACGT TTAACAGGTA GTTTATGGAG GTTTGTTGAA
115681  TGAAAGGATG TGTGAATTTT CTATGTAAGT CTCCAGGCTC TCCACTAAGC CCACCAGAAT
115741  GCTAACACAA TCAATTCCCC ATCTCATTCC TTGACCTGCC ACTGCCTGAA GCAATCAGCG
115801  TGCAGTTTCT CTTTAGAAAA TCTGGGGGAT AGTCTAGGGG TTGCAAATTA AGCAACATTA
115861  TCTTTGTTCT GAACAAGGAC TGCATGAGTG TTAGGACTGA AGAAGGCCCA AGGTGGTGGT
115921  GGGTATGCCT AAGATGAGTA TGACATATCA GCAATGCTAT GAACATAGCA ATGCTATGAA
115981  AGGCCAGGCA AAACGTAACA GGAGCTAGTC GTGGCTTATT GTTACAACGA CTATACCTCC
116041  CATATGGGTA ATCGATATCC ACACACCCCT CTACATTGAC TCTGGAATTC AGGAAAGGGA
116101  ATTAAAATTT TCTAACTTAT GTACCCAAT GATTTCAACA ATATCTGGCA TATGAGATCA
116161  ATAAATATCT TTAAATACC AACTAAGAAA GACATAAAAT GACCCACCCT CCATACCAGG
116221  CTCATTTTTG CTCCTCTGAT TCCTGAAACT ATCCAGAATG CAGCTATGAA TTCTCTCCAT
116281  TGTCAGTTTT AAATTAAGCC AAGCTGGGTA CTTGTGTAAT TCCTCAAGAA ATCCTGGATG
116341  AAAACTGTCA GGTGGAAAAC AGGACCTCAA AATAAAGAGA CATCCATCAC TGAAGCTAAC
116401  ATCGTGAGGC TGAAATCAGT CCTATAACAA TGGTACCAAA AAGAGCACAA TGAGAGGCAT
116461  TTGTGAATAT TTACTCAGAT GAGAGTAAGA TATTTCCCTA TCAGCTAACC TGAAGTTCAC
116521  ATCCCTTTTC CAGCTGAGTT CTGAAGCTAG ATGTACTTAA CTGGAACACA TAACTGCATC
116581  AGGAACATCC TTTAAAACTA TGGCTACAAT GGCTTGACTG GACAAACCCC AGGCTTCCAG
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116641    GTTTAGCACA GGTGGCCCTT CACAGACCAA CATTGCCTAT GCTACCAACC TCATGTCCTA
116701    CCACCCTGCT TGCATCATTT CTCTCTCTGC ATATATAAAA ATATATGTGT ATGTATATAA
116761    TCAGCTTTAT TGATATTTAA TATACCACAA AATTTGCCCA CTTTAGGTAC AGTTCAATGA
116821    ATTTTACCGT GTTTTCTTAG TTGTACAACC ATCATCACAA TTTAATTTCG GAATATTTCT
116881    ATCACCCAAA TTTCCATTTC TGCGTAAAGG GGGAAAAAAA AAGGTTAACT GCTGAAGGCC
116941    GCGGTAACAC TGAAAAAGGT GCCTTTTCTC TCTAAACAG ATTTTAATCT CCCCTGAATT
117001    TAGTGTCCTG GGTATTCCAG GAGTCTGAAT AGGGTTTCAA TTTTCAGGGT CTTTTTAATA
117061    GAGTAAAACT GTATTGGTGG CGATAAATTT AGTATTGCTC TCAGTACATG ATTGAGGGAT
117121    ACTTAAATGT CTCTGTGATT TTATTTCATA ATCGCTAAAA GATGGTTTTT TTTTTTCCTA
117181    AAACAGGGTT TTTGTTTTTT CTCAATAAGC TTCTTAGCTT CCCCTCCGGC TCCCTGGCTT
117241    GCCTCAGGAA ATATTAGCTC ATCAGTTCTG ATTGGTTGAC AGCTACGAAT GGCCCTCATT
117301    GATTGGGCAG CGCTTCTTTG TCCCTTGGAA ACTAATACAA ATTTTTAACA CTACTTTTTT
117361    TCCACTCTTT CTTCAGAGTT GGAATATCGT TGCTCCCCTA CCCATATGTA GTGAGTGGAG
117421    GGCAAACTTG GAGTTCCCCT AATCTTTCCT TTTTAGGATG TCAGCTCAGT ATCATTCATC
117481    TTAATTACAC ATTGAGCTTC TTGACTTAAT GGATACAGCT CTTCTTTTGT TTAGTTGGGC
117541    GGCCCTGAAA AGGGCCTTTG GTTCAGAAAT GCAAGCTGTG GAGAAATCAG CAACCTTAAC
117601    CGCCAAAGCC ATAAAGGGTG CGTCCCTGGC GCTTAAGCGC GTAGACCACG TCCATGGCAG
117661    TGACTGTCTT GCGCTTGGCG TGCTCCGTAT AGGTGACAGC GTCACGGATC ACGTTCTCCA
117721    AAAACACCTT GAGCACCCCG CGAGTCTCCT CGTAGATCAG ACCAGAGATC CGCTTCACAC
117781    CGCCACGCCG GGCCAGACGC CGGATGGCCG GCTTGGTGAT GCCCTGGATG TTGTCACGCA
117841    ACACCTTGCG GTGGCGCTTG GCACCCCCCT TACCCAAACC CTTCCCGCCC TTACCACGTC
117901    CAGACATGAC TTCCCAAGAA GTGAACCAAG AGCAAGTGAG AGAATAGGAA ACCGATCTTT
117961    ATATATCTAC GTTACCCCTG CCCCCACCTC CAGCGGACAC AGAGACTGAA AAGCGCGCAG
118021    GCGGGAAATG TGACGCCTAC AGTCCGCTCC TTTAACCCCT CCTCCAAGCC CCAGGAAATG
118081    GCGGGAGCAG CGATTGGGGG AGGGTGGGGA GATGAGGGTG GGACCAAGCA GGCTTGACCA
118141    ATGGCCTTTA TTTTCTTAAC AGAGCTACAG GCTTTGAGGA ACTGGGTTAA GAATTAAATG
118201    TAAACCCATT CTGACTCCAG AATTATTTTA AGTCGAACTT TTTTTTTAAC CGAATCTCTC
118261    TGTCGCCCAG ACTGGAGTAC ATTAGAGCCA TCTCGATTCA CTGAAACCTC TGCCTCTCAG
118321    GTTCAAGTGT TTCTCCTGCC TCAGCCTTCA GAGTGTACCT GGGATTACAA GCGCTCGCCG
118381    TCGCGCCCGG CGTGTTTTTG TATTTTTCGT AGAGACGGGA TTCGGCCATG TTGGCCAGGC
118441    TGATCCCGAA CTCCTGATTT CTGGTAATCC GCCCGCCTCA GCCTCTTAAA GTGCTTGAAT
118501    TACAGGCGTG AGTCACCGCG ACCGGCCGAA ATCGATTGGT TTTGAAGCCT TCAGTAGCAT
118561    TAAAACGAAA AGTGCTCCCA ATGCATTCCC TTTTGTCTTA AATTGGTTTC TTACAGCTAC
118621    TTTACTTGAA AAGGTGGTGG CTCTGAAAAG AGCCTTTGCT TGGACCGTCA GAGAGACCAC
118681    AGTAATCACG CCCTCTCTCC GCGGATGCGG CGGGCGAGCT GGATGTCCTT GGGCATGATA
118741    GTGACGCGCT TGGCGTGGAT GGCGCACAGG TTAGTGTCCT CAAATAGCCC TACCAAGTAG
118801    GCCTCGCACG CCTCCTGCAG AGCCATCACA GCGGAGCTCT GGAAACGCAG GTCTGTTTTA
118861    AAGTCCTGCG CAATCTCGCG CACCAGGCGC TGGAAAGGTA GTTTACGAAT AAGCAGTTCA
118921    GTGGACTTCT GATAACGGCG GATCTCGCGC AGAGCCACGG TGCCCGGCCG GTAGCGGTGG
118981    GGCTTTTTCA CGCCGCCGGT GGCCGGAGCG CTTTTGCGGG CTGCCTTAGT GGCCAACTGT
119041    TTGCGTGGCG CCTTGCCACC AGTAGACTTC CGAGCAGTTT GCTTAGTGCG AGCCATGACG
119101    GAAAAACAGC ACAGCGGAAC ACCCAACACT AGCGCAAATA CGCCCATGAG CTGCTCTATT
119161    TATAGTGTGT AAAGTGCAGT GATTGGATGA TAGAAGACGC TAAATATGAC GTTACACACT
119221    CTGATTGGTC TATCTTTAAG CCAGCAACAA TCGTGCAGTT TCACCGGCTA CTATATTCTA
119281    TTCCAACTCT ACAGATGATT ATTTAAGTGG TATTTTATTA CTACTATTAT TTTATTTTAC
119341    TTTTGCTTTG TTCCCCAAGC TGGTCTTAAA CTTGGGCTCA AAAGATCTTC CCGCCTCAGC
119401    ATCCAGAGTA GCTGGGATTA CAGGGGAGCC CCACTGCGCC GGCTTGGACT TTAATTTTTT
119461    AAACTTGTCC TCTTCTACAT CTGGTTTTCA TAACCTGAAG CTGTGTTTA TTTTCCATAA
119521    AACAAGGCAT TGATTCCAAA GGTATTATAA TTCCCCAATT CCGTATAACC TTCAGCTCTT
119581    TAGGAAAAAA AAAAAAAAA AAAAAGAGG GAATACTGCT CACCTCCTCT CCGGAAATGT
119641    ACCCTTTACG GGAATTTCTG AAACCTTTCA CAAGAATTGG ATTCCTTTGT AATGCTTTAA
119701    TTGACTTAGG AGTGTTATTG AAATCTACAA AGCATCTCAA ACATAGTAGG ATTACACTAT
119761    TACTCAGAAA CATTTTCTAT GAGACGTCTT TCTCTTGATT ATGCTCTTTG AATCCTAAAC
119821    TTGCAGCGTT CTGCAGCTTT TGTTTTCTAA AGCCTAGGTG TACTCTGCCA GTCACAAAAT
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119881   GGCGTTTCTC CAGCACTGCC GCCAGGTACC ACCAGCTGGG AGTTGTTCCT CTTGCGGAGC
119941   AGGAGGTGGA CTTGGCCCAA GAGAAACTGG ATAGTGGTTC GCAAGGAACA TAATTTAGCA
120001   TTGCCAAGAG CTAATGCAAT CATTTGAAA ATCTCAAAAC ACTGAAAAGT GGATTGTGAC
120061   CTTTTTAAAT TCACAAGAGA CAGGCCACAT TCTATCTTTT GATTGGTTTA GGCTATTTTC
120121   TTGAACAGCC ATTTAGAAAG CAGATCTATC ATCCTTCATT TGCATGGAGC GTTCCCATTT
120181   TATTTGAAAC CAGTTTAACC CAATAGAAAA AAGGGAGGCA GAACCCATTA TTTAAAGTGG
120241   AAACTCCTGA ATCAGATAAT TAGGAGTATT TCCTTTTCAA AAGTTGCGTT TTTTCAGATA
120301   CCTCGCTTAT TACACTAAGA AAGGTTTATA TCTTTCACAA AGGGTTTACT TACAAAAATC
120361   TTCCAATTTT GTATACCTGT GTTTCATAAC TGACTAGCCG TCAAACCAAG ATGTAGAGTT
120421   TCCAACCGTT ATTTTCCAAA TTTTTAGAAA TTACGTGAAA TATTTGAATG CATGCCTTCT
120481   CAATAAAATG GGACGTAGGA AGCACTGGTG CAGAAGATGG GTACAATACT TATCTGGGAC
120541   CACTCCATTA TTTGGTTGGC ACGTTGTTTG AAGAAAAAGG GGAAAAGCTC AGGTTACTTA
120601   GCATGGTTCG GACTTATTTG AAAACTACCA CAGCAGGAGC GGAAATAAGA CCGCATTACC
120661   TCACTCTCTG CTGTGCTGTG CTAGGGGGTT ATCCAGAATA GGATTGTAGA AGTGGATGTC
120721   GATTAATAG TTTTTTATTC TCCCATTAGC TGAGTCTCTG ATTGGCAATG TGAGATCGTT
120781   TTAGCTTATT GATACTTTGA AATGCACTTA ACAGCCACAA ACAAGTTAAA GGGTTGTTAC
120841   CATAAAATCT TATCCCCAGG GTGTGCTTGC ATTTATCACC CGTGTTTGCT TTCACACTAA
120901   GTGGACTTAA CTCCCCAGCA GAATGCCTGT CAGGGAACCG GTTTCGTGGA CCCAGCATTT
120961   AACGCCTTTC GCAGGCTTGT GAGGCCCATA AATATTTGTT GAATAAAAGA ATGAGTTGAC
121021   CATGTCATGG TGCGCTGATT GCGTGTGCTG ACATGGAACA CAGGTTGTAA ACCTTAATAC
121081   CAATTTGGGG CATGTTGTAT GGATGAAAAG GGCATTGGAA ATTCCTGAAG TGCATCCCAC
121141   ATTGGACTGT GGAAATAAGT TGCAAGTGCA GAAACGTTTC CACACTTGCA GTTTGAGTAT
121201   TAATTGCAGC GTTTGTGAAT TCTGGTGTTG TCTACGATTC ATTCTTGTTT GACGTGAAAG
121261   GTATTCGCGA GACACATCGC TCTAAAACAT TGCCAGAAAA TGTAATAGAG TTGATGACAA
121321   CTGGCCCTAA CACGGCCTAA AACTCGCACT TTTCTCTCCC TCCGCAACTA TTCAAAACAC
121381   TGTATTTTAC ATTTCTTGCA AATTAAAAAC TAACATCTCT GGCAACGGAC CTCTAAAAAT
121441   TTCTAATAAA ACTCCTCGGA TGCTTGTGGC ACTGCATTTG TAAACCGCCC CCTCTCAACC
121501   TACTCCCTAA AAAAGAGCTG CTTTTTGAGA GAGAAGCGGT ACCCTCTGAT GTTACTGGGC
121561   GGCAGTCTGC CTACAATTTC CTTCACAATG AGGCAACCAG AGCGGCTTTT TCTGTGTGTT
121621   TGCTTGCGTT GAGGGGAGCA GGACCATAGG CCCTAGAGGC CCCCAGCTGC CTTCTGAGAC
121681   TGGGCGAAAC CCTCGGCAGC GCGCAGGGGG CGCTAGGGCG CGAGGGGCGG GCACTGACGG
121741   GCACCAATCA CGGCGCAGTC CCACCCTATA AATAGGCTGC GTTGGGGCCT TTTTTTCGCA
121801   TCCTGCTTCG TCAGGTTTAT ACCACTTTAT TTGGTGTGCT GTGTTAGTCA CCATGTCTGA
121861   AACAGTGCCT CCCGCCCCCG CCGCTTCTGC TGCTCCTGAG AAACCTTTAG CTGGCAAGAA
121921   GGCAAAGAAA CCTGCTAAGG CTGCAGCAGC CTCCAAGAAA AAACCCGCTG GCCCTTCCGT
121981   GTCAGAGCTG ATCGTGCAGG CTGCTTCCTC CTCTAAGGAG CGTGGTGGTG TGTCGTTGGC
122041   AGCTCTTAAA AAGGCGCTGG CGGCCGCAGG CTACGACGTG GAGAAGAACA ACAGCCGCAT
122101   TAAGCTGGGC ATTAAGAGCC TGGTAAGCAA GGGAACGTTG GTGCAGACAA AGGGTACCGG
122161   AGCCTCGGGT TCCTTCAAGC TCAACAAGAA GGCGTCCTCC GTGGAAACCA AGCCCGGCGC
122221   CTCAAAGGTG GCTACAAAAA CTAAGGCAAC GGGTGCATCT AAAAAGCTCA AAAAGGCCAC
122281   GGGGGCTAGC AAAAAGAGCG TCAAGACTCC GAAAAAGGCT AAAAAGCCTG CGGCAACAAG
122341   GAAATCCTCC AAGAATCCAA AAAAACCCAA AACTGTAAAG CCCAAGAAAG TAGCTAAAAG
122401   CCCTGCTAAA GCTAAGGCTG TAAAACCCAA GGCGGCCAAG GCTAGGGTGA CGAAGCCAAA
122461   GACTGCCAAA CCCAAGAAAG CGGCACCCAA GAAAAAGTAA ATTCAGTTAG AAGTTTCTTC
122521   TAGTAACCCA ACGGCTCTTT TAAGAGCCAC CTACGCATTT CAGGAAAAGA GCTGTAGTAC
122581   ACAGATGAAA TCCCCCAAGC AAATGCAACA CGCCCTCAAT TATATTAGAA TCACTTGGAG
122641   AGTCGATAGA ACTTTAACAT AGCCTCATCT AGTAAGAATT TACTACTCAA TCTATCAAAG
122701   ATAGCAAGGT GAATTCAAAT GCACCGAGTT AAAATCGAGT TTTAAAGTCA CCTGGGTTTC
122761   GGTAGCCGGA AGTCCGCGT CTCACGACTC CAAGCTAATT AGTCATAACC GTATTGAACC
122821   AAGGTTGAAG CCCAGTCCCA GGCTTGAGGC TTTTTATTAT ACAAGGTTAA AGTGGGGATA
122881   TTGCGTTTTG GGGTCAATAT TGCTAAAGTA GCATTTTCCG AAATTGGGTG GTCCTAAGAA
122941   ATGCTTCTGG GATAGTTGGC AAAAATATATG GCTTAACCAC GCCCTCTCCA CAGGAGTGGC
123001   TAGCGAGCTG TCTGTCCTTG GGAAGGACGG TGACCCTGCT GGCGTGGCTG CGCCCCACGT
123061   TGGCGTCCTC TGAAAGCCCC GCCAGGTAGG CCTAGCTCGC TTGCTTTCTG CAGCGCCATC
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123121    ATGACAAAGC TTTGAAACGC AAAATGCTTT CTTTGTGCAG CGCCTTACCA TGGGTGCACT
123181    TACGGGCTGT CGACTTGGTT TAGGCCCTTG TCAGGACAAA GGAGCTTAGT TTGTTGGAGT
123241    TTTAGAGCTG CAACCCAAAA TCCCTTGCTC GGTTTCTCTG TTTTTAGAAA CGGAAGCGCC
123301    CTGATTGGAT ATTTGAAAAT TACTGTGCTT AACTGGATCG TGTTTCATCA ATCGTGCAGG
123361    ATTTTCAACC CTGGTGGAGC CCACACATTC AAAACTGAAG ATCCTTTTCT CAGAACTGCC
123421    CCTTTAAGCT TTTGCAATTT TAATTCTGGG GGTCAGATTT TAATAATTGG ACTTTTTTGT
123481    TTACATCTGA CAAGAGTATA TGATGAGCCA AGTTTACTCA CTTTTACTTA GTGCAGTTCA
123541    ATTCTAAAAG TTTATTTTTG CGTGTGTGCA TATGAGTTAA TAATCAGTTG TATTTTTCAA
123601    ACGGTCTTTT TTCAATTGTT TTGCTTAGCT CCTTCCATCG TCTAAAGTCA GGGATACAGG
123661    CACATCACAT CCCTGTTCCC CCTTCCTCAA ACTAATATGT AGCTACCTAG GTTTATCCTT
123721    TAAAACAAAA ATTCTCACCT ATTTTTGTGA GAAATATACA TGTTTTTCTT TGAACTAAGT
123781    ATTTTACATA CACCTATCTA TATACATGCA TACTTGTGGT TTTGTTTTTT TAAAAAAAAA
123841    AAAAAAAAAA CACGTTATCT TTTGAGACTG GGTCTCAGTC TGTTGCCCAG ACTGGACTGC
123901    AGTGGCATAA TCACAGCACA CTGTAACCTC CAACTCCTGG GCTCAGGCTA TCCTGCAGCC
123961    TCAGCATCCG GAGTAGCTGG GATTGCATGC ACGCACCACC AAGCCGGGCT TTTTGTTTTT
124021    ATTTTTTGTG GAGACAGTCA CACCATGTTG TCCAAGCTGG TCTAGAAATG GCCTCAAGTG
124081    ATCATCGACC TCCCAAAGTG TTGGGATTAC GGTCACTGTG CCTGGCCTTG TATGCATAAT
124141    TGTTTTGTCT TTTGATTAGG GTTATTAATT TAAAAAACAA AGCCTGGACG CAGTGGCTCA
124201    CATCTGTAAT CCCAGCACTT TAGGAAGCCG GATGGGCAGA TTACTTGAGC TCAGGAGTTC
124261    AAGACCAGCC TGGGCAACAT GGTGAAATCC CATCTTGACA AAAAATACAA AAAATTAGCA
124321    AGGCCCAGTG GCACGCACTT ATAGTCCCAG CTACTTGGGA GGCTGGGGTG GGAAGATGAC
124381    TGGAACCTGG GAGGTAGAGG CTGCAGTGAG CAGAGATCGT GCCACTGCAC TCAAGCCTAG
124441    GTGACAGAAT GAGACCCAGT CTCAAAACAA AAATAATAAA AATTTTTTAC AACGATGTTA
124501    TATACACTTC TGCATGTTGC TTTTCTCTTA ACCAAACTTT TCTAAAACCC TGTCATGAAA
124561    AAAGAAATCC TTCACATGGA ATAGCATAAG TTATTCATCC ATTTCTTATT GATAAGCATT
124621    GATGTTTCCA GTTACCACTG CTGAACATGG TGCAATTGAA TAGAATTCCA GGGCTGAGAT
124681    TGCTAGGTTT TAGGTTGTAT TTTATTATTT TATTTATTTA TTTATTTATT TAGACAGAGT
124741    CTTACTCTGT CACCCATGGT GGAGTACAGT GCCATGACCT CAGTTGCAAC CTTTGCCTCC
124801    TGAGTTCAAG CGATTCTCAT GCCTCCGGTC TCCCGAGTAG CTGGGATTAC AGGCACCTGC
124861    CACCAGGCCT GGCTAATTTT TGTATTTTTA GGAGAGATGG GGTTTCACCA TGTTGGCCAG
124921    ACTGGTCTCA AACTCCTGGC CTCAAGTGAT CTGGCCACCT CGGCCTCCCG AAGTGCTGGG
124981    ATTACAGGTG TGAGCCATGG CTCCAGACCT GGACTTTGTC TTCTGTTTCA TCAGTCCTTC
125041    TGTTGGTTCA AGCACAGTAT CACACTGAAG ACTGATGATT CTATATAAAT ATGGTAAAGA
125101    CTGTACACCC TAACTGTTCT TATTTTTTAA TTTTAAGGCA ATTTTAGATT CCAGCTTTCC
125161    AAAGAATTGT GGAATGCTTA GAGCTAGAGA AGCCTTGGAA GTCATTTAGT TTTTGTTTTG
125221    TCAGAGAAAA TTCTGTAGAG ACTCTGTCCT GCTCTCACTG AATACCATCC CATAGTACCC
125281    CCCAACAGCT TTAAAGGGCA ATAATACCTT ATGGACAGTA TGCTTTTCCT CAAATATATT
125341    CTAAGCCATG GTCAATGCAA AAGAGTGAGA AGGAAAGTAG AATAAGTTAT CTAAGAATCA
125401    GTGGGTGCTC TCTTTAAACT GATTTATCAC TCCCCCTTCC AAACTCTCTT GAAGGTCACT
125461    CTGCCTCCCT TTCTACATAA GAACTCCTAA CTCCAAGGGA GGAAGGTAAG TTATTCTTAT
125521    TCCTTGCTTA GAAAAAGAGA AAATAGGTTT GGTAAGCATC CGCTTTCTGC TACCATTCTC
125581    TGTGTTTCTG TGTTTTTTAT AGGATCATTC AATTATTGGT TGGCTCTTGA GAGGGAATGC
125641    AAGGTTCAAG GACACAAGCC TAGATCTTGC CTGTATAGAA CCTCATGATG TTATGCTTCT
125701    CTAAAATGAG GCCTGGAGGA GACATGTTGA AAGTGACCCA TAAATCTGCA GTATCTCATG
125761    TCTCTCAATG GGGACAAGGA GTACCATGGG AAATAGCATT AGGTCAATGA CAGTAACAAC
125821    TCCCAGGTGA GTTGATTTAT TCTTTTATTT ATAAAGTTGT TAATATGCTA CATAGTCCCT
125881    AATTTTGCCA CAAATAGTCA TTATTTTAAT TTCATATTTC ACTATTGATA AATGAAGGAA
125941    AAAATGAGTA GCAGTTAAGC AGTCCATAAA CCTACATATA AAGCAAATTG GAGATTTTAA
126001    AATTGATTCT GGATGCTTAA AATCCTTCTC ATTGAAAAAA AATTTCGTAT TAGAAGATTT
126061    CAACATTCTT TAAACTGAGA AGCATAACAT ATAAACAGAA AACCACAGCA AAACAAAAAT
126121    GCAAAGCTCA ATAAATGAAC ACAAAGTGAA CACCATAATA ATTGCCACAC AAGTAAAAAA
126181    ACAGAAAATC AGCCAACCCT CCCAGAGCTG CCTGATGCTT GCTTCCAGTC ACATTATCAC
126241    TCCATCTGCC CTAAACATAA CCCCTATTTT GATTTCCAAT GCTGTAATTT AGTATGCCTG
126301    TTTTTGAAAC ATATAAAATG GAAATAAAAC AAATGTAATC CTATGTACCT GACATATTTC
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126361   ACTCCAGAAC ATTAGGTTTG AATAGATTCA TCTGTGTTGC TGTGTATAAC TTTAATTCAT
126421   TTTTATTGTT ATGTAATATT CCATGTTATG AGTGCAACAA TTTAGGTGTC TACTGTTGAT
126481   GCATATTTGC TTCCCTTTTT CAGCTAATAT AAACAATACC GTGAATATTC CTGTGTATGT
126541   GTCTTGGTAT ATATAGGAAT ACATATTTTG TTTGTATACC TAGGAGAGGA ATTGTTGGGT
126601   CAAATGCTAA ACTCTTTTTG AAAGTGGTGA TATTAGGTTT ACATGCGATG AAATGAAAAT
126661   TAAAACCACA GTTATAAACA GCATGGATGA ACCTCACAAA CCTAATGTTG ATGGAATCTA
126721   GCTGGGAATT CCTGTTCTTC CATATACTTC CCAATATTTT TTTCCAATTA AAATTGTTAA
126781   TCTTTTGAAG ATGTTATCCA TTGTGGCAGA GTGTGCAGTAT TATCTCATTA TGGTTTTATT
126841   TTACATCTTT TGCCCATTTT TTCTTAATTG GATTGTATAT CAGTCGACTT GGGCTGCCAT
126901   AACAAAAATA CTAGACTAGG TAGCTTGAAC AAAAGGAATT TATTACCTCA CAGTTCTAAA
126961   GGCCAGGCCA GAAATCCTAA ATTGAGGTGC CAAGAGATTC AGTTTCTAGT GAGGGCTCTC
127021   TTATTGACCT GAAGATAGTT GCTGTCTTAG ATTGTTTGGT GCTGAACAGA ATACCAGAGA
127081   CCAAATAATT TATAAAGAAT ACAGATTTAT TTCTTACAAT TCTGGTGGCT ATAAAGCCTA
127141   TGGTCGAGGG GCCCACCTCT GGCAAGGGCC TTCTTACTGT TATGGCAGAT GTGAGATGTC
127201   ATCTCATATT CAAACCACAG CAGTCGCCTT TTGTGTCCTC ATGTGGCCTC TTCATATGCC
127261   CATAAAATGA CCTCATGTCT CTTCCTTTTC TTATAAGGAC ACCAGATCTA TCAGACTACT
127321   GGCCTACTCT TATGACCTCA TTTAACCTTA AATATCTCCA TAAAGTCCCA AAATCCCTAT
127381   CTCCAAATAT AGGCACATTG GGTGTTAGAG TTTCAACATC AATTTTGGGG GAACACAATT
127441   TAGGCCAAAA AGATTGTGTT TTTTCTTGTT GGTTTAAGAT AGCTGTCTTT TTGTCCTTTT
127501   TGTCCTTTCT TTTTTTTTGA GGTGGACTCT TGCTGTGTCA CCCGGGTTGG AGTGCAGTGG
127561   CGCTGTCTCA GCTCACTGCA ACCTCCACCT CCTGGGTTCA AGAAATTCTC CTCCTCCCAA
127621   GTAGCTGGGA CTACAGGTGC ATACCACCGC GCCCTGCTAA TTTTTGTATT TTTGATAGAG
127681   ACGGGGTTTC ACCATGTTGG CCAGGCTGGT CTCAAACTCC TGACCTCAGG TGATCCACCT
127741   GCCTCGGCCT CCCAAAATGC TGAGATTACA GGTGTGAGCC ACCAAACCTG GCCTGTCTTT
127801   TCTGTTTTAA GTTTTTAAAT TTTGCTCACG AACCCTTTAT CCATTTTATG TGTTGCAGGT
127861   ATTTCCTCTG TAACTTGTCT TCACTCTGTC AGAGGCTGGA GTGCAGTGGC ACAATCACAG
127921   CTCACTGCAG CCTCCACCTC CCAGGATCAA GCGATCCTCC CATCTTATCC TCCTTAGTAG
127981   GTGGGACTAC ATGTGCAGGC CACCATGCCC AGCTAATCTT TGTATTTTTT TGTAGAGATG
128041   GTGCTGTTGC CCAAGTTGGT CTCAAACTCC TGAGCTCAAG CAATCCATCA ACCTTGGCCT
128101   CCCAAAGTGT TGGGACTAGA GGTGTGAGCC ACCACTGCAC CCAGCCAATG ATATCTCATG
128161   ATGCATTAAA GTCATTAATT TAGTGTACTC AAATTAAGCA CACTGCCCTT TTATGCACAA
128221   CCTTTTTTGT ATCTTATTTA AAAAATCATT TTCTATTTCA AGGTCATGAA GATCTTATTT
128281   TATAATACCT TCTTGTGAAA TTAGTTCTCA AGACTACCCT CACTTCTAAC ACCAATTATA
128341   AGTTGGGAGG TCTGTGGTTC CCAATCAACC TTAGGTTAGT AATTTGCTAA AAGGACTCAC
128401   AGAACTTGCT GAAGCTGTTA GCCTCATGGT TACAATTTAT TATAGGATAT ATAGCTTATT
128461   ATGTCATTCC AATGCAATGT AAAATTATAC AACTACTTTT AAAAAGATTT TAGCATTTGA
128521   CCCAACAATT TCACTCTGAG GTATACAAAC AGCAGATATG TGTGCACATA TATACCAAGA
128581   CACATACACA GCAAAATTCA TTGTTTGTAA TAGTTGAAAA GGGGAAACAA CTCAAGGAAT
128641   AAAGATTAAA ATCAGCTGAG AAAAGAAACA CACAAGGCAG TATTATGGAT CGAATTGTAT
128701   GCAGATCTCC CTTGCCCCCA GAAGATATGT TTAAAGTCCC AACTCCCAGT ACCTCAGAAT
128761   TGTGGCCTTA TTTGGAAATA GGATAGTTGC AGATATAATT AGTTAAGATG AGGTTATAGT
128821   ACAGTATGAT GGGCTGGTGA CTTAGAAGAA GTAGTATATA TATATTTTTT AATAGAACTA
128881   GTATTCTTCT AAGGTGGTCA CGTGAAGACA GACACACACA GGCAGAGACT GCGGTTATGC
128941   AGCTGCAGGT CAAGGAATGT CAAAGGTTGC CAGCAAGTAC GAGAAGCTAG GAAGAGTCAA
129001   GGAAGGATTT TCCTACAGGC TTCAGTGGAA GCATAGATCT AATGATACCT TCATGTCAGA
129061   TTTCTAGCTT CCAGAACTAC AAGAGAATAT ATTTGTTGTT TTAAGCCACC CTAGCTTCTA
129121   GCTCTTTGTT ACAGCAGCCC TAGGAAACTA ATATAGGCAC AATCCAGGCA AGTTCCAAAT
129181   ATGAGCTTCC AGTTGTCCTC TCCCAGTAAT ATGAACAGTA TTACTTTCCC AGCATTAATG
129241   TGTGACAATA CACATGACGT ACAGAGCAGT CCCCACTTAT GCACAAAACA TATGTTCCAG
129301   GACCTCCAGT GGATGTCTGA AACCATGGAT AGTACTGAAC TCTATATAGC TGTTTTTTCC
129361   TATACAGACA CAGCTATGAT AAGGCTTAAT TTATAAATTA GGCACAGTAA GAGATTAATA
129421   ACAATAAATT AGAATAATTG TTAAGAATAT ACTGTATAAA AGTTAGGTGA ATGTTTATTT
129481   CTGAAATTTA CCGTTTATTA TTTTTGGACT GCAGTAGACC ACAGGAACTA AAACCATGTA
129541   GAAACCGTAT ACAAGAGAAC TGTATTTCAC CCGAGCCTCA GTGTGCAGTT TTAATGGCCT
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129601   GCCATGGTTG ACTGCTCACA TGGCCGATCT TTTAGTCTAC CTCCACAGGT AGAGCTGATA
129661   CTGTGTGGCT CAAAGTTCCT ATTATAAATC ACATTGTTGA CTGTGTGGTG GTCAAAACCT
129721   CCAGGTAAAC AAAGACACAC TTATCAGTGA GAACATTTCA AGGGTCTAAA ATTCATCTCC
129781   CAGTAGCTGA GGGCAAAGGC TAGACCTCTT TTTGGGTAAG ATAAATTTTT TACCATATAC
129841   TTTATTTTGC TTTTCATGTT TAACTTTATT TTGCTTTTCA TGTTAGTTCC CCTGGAATTG
129901   TTTTTTGTGT ATAGTGTGAA GTAGGGGGTC AAGTTTCTTT TTTTTTCCTT TTTGTTCTTT
129961   TTCTGTTTAA AAGGCTATAC AATTGTCCCA TGCCATTTAT TTACAAGAGT CCTTTCACCA
130021   TTGTTGTATG GTGCCACTTT AGATGTAAAT CAATGTCCAT ATTTGTTTGA GCCTGTTCCA
130081   TTCGTTTGTC TATTTTTGGA CAACACTGCC CTGATTATTG TCATTTTATC AGTTTTGATA
130141   TTTAATAAAG CAACAGATTT GTTTATTTTG GGCCCTTGGA TTTGTGTATT AAATTTGAAC
130201   CCTGTTTGTC AATTTCTATA ATAAAGCTTA TTGGGAATCT GATTAGGATT ACAATGGTTT
130261   TGTAGATCAG TTTGGGGACA ATTAATACCT TTAAAATATT GACCGCTTCA ACTGTAAATA
130321   TACTCCTCCA TTATTTAGTT TTCCTGTTTA ATTTATCTGA GTAATACATT ATAGTTTTCT
130381   TCGTAGAAGT CAGATACGTA GAAAATTCAA AGCCCAAGTG CAATAGCTCA TGTCTGTAAT
130441   ACCAGCACTT TGGGAGGCCG ATGTGGGTGG ATCACCTGAG GTCAGGAGTT TGAGACCAGA
130501   CTGGCCAACA TGGTGAAACC TCATCTCTAG TAAAAATACA AAAATTAGCT GGGTGTGGTG
130561   GCGGGCACCT GTAATCCCAG CTAATCAGGA GACTGAGGCA GGAGAATCGC TTGAACCCAG
130621   GAGGCAGAGG TTGCAGTGAG CCAAGTTCCT GTCACTGCAC CCCACCCTGG GCGACAGAGC
130681   GAGACTTCGT CTCAAAAAAA CAAAAAAAAG AACATTCAAA TAATCAATGT AGATAATTCA
130741   AATAACTAAA AAATGAACAG TTATTAAAAT ATCAGGATAT AAAAGCAAAA AAATCAATAA
130801   CCTCCATATA TACAAAATGG CCAGTTAGAG AAAAAAAAAA GAATAGGCGA GACTTAAAAA
130861   GGCTGGGAAT CTCCCTGAAA ATCTTTGAGA GCCTTGGCCC TGCCCTCAGG GATTTCTCTG
130921   GCTTCATGCC CAGATATGGG TACAGTTCCT TGTTTAAAAA AATTTTGCTC CATCAATCAA
130981   CAAGGGGCTC CTTCCTCAGA GCACAAGGAC CTCCATAACA CCGGACACTA GATGTCTAAG
131041   GGACACCTCT TAAGGAAGTT AGACTTCCAA AGAATGGTGT TTCCTCTGTC CCCAAACTCT
131101   GGAACTCACA GCACAACTGC TCCTTGGAGT TCGGTTTCAA ATCTACAAGG CTGTCATGGA
131161   GGTTGCAGAC CAAGTCCGTG GCCTCAGTGT CCGGATGTAC GGTGGCCTTG GCACCTGAAT
131221   GTGAGAACAT GACCTCCCTG AAACCACCAC AAGTATTGTT TCATGTTATG TATGTTTTTT
131281   CTTATCTGAA ATTCCTTTTC TTTAAAAATT CAAATTACAT ATTTTTCAAG CCCCTGAACA
131341   AGCTTCATGA GCATTTATTG AACCCACAGC TTTTAAAACC TACTGAACAC TTTGCTCTAT
131401   GTTGTCATTC ACTATCCACC AATTATTTAA TTATTGATCA ATATTGTTTC CTTAGTGTTG
131461   GGATCATTTA TGCATGTATT TCTTTTATAT TGCATATTTT ATATTTCTGC ATTACAGTTA
131521   TTACATATTA CTTTTGCTAC AGTAATAGTT CAGAAGTGTA CATCCAAAAT TTAGCTGTGA
131581   AGTGGATGGA CTGAGGCAGA ACTGGAGGCA AGAAAATGTC ACAGTAATTC TAAAAAAGAT
131641   GATGTACAAT TAGAGCAAGA GAGTAGCACT GAAATTGAAG AAAAATAGAT GCGTTTGAGA
131701   GAAAATTAGG AGGTAGAATC AACAGATTAG ATGTAGGGAT GAGAAGGGTC AAAGATGACA
131761   CTAGGGTTTT TAACTGGAGC AAGTAGGTAG ACAGAACATT TCTTCCTGAA AGGGCAGGTC
131821   AGATCATGTG TTGTCTCAAA GGGCATGAAG AGTAGAAAGC CTGGGACAGA TCCTGAGATG
131881   ACCAATACCC ATGGTGCAGG GAGAGGGAGG GAGATCTGCT AAAAAGACTG CAAATGTCAG
131941   GATAGTAGAA AATCATGAGT GTGTGATGTC CTGGAAGTTG AGACAGTATC ACATTTGAGA
132001   ACATTTAAAT TGGTAACTCT GACAAAACCT GGAGGCCAAC TGTGAATGCC CATGAGAGTG
132061   AGAAGCTCCC ACACTTTTGT GGGCATCAGA AAGCCCACCA GGTTCCTGCA GTGAAGATCT
132121   GAGAAGGATC CTCTTGTGGC TTTGGCAGGG AGAGAAGAAT TATTATGAAA TACACCCCAG
132181   AACCTTCTTC AAAACAAAGG CCTACTCTCA AGGGGAAAAC ATTTTGCCAG AGTCTTATCC
132241   CAGCTGGGAG AAGGTAATTC TTCCCACTGC AGCCTCATCT AGGCTTTCTG TCTCACTTAA
132301   GGGAAGAAAA TTAGTCAACA GGGATCAGAG CTTCATGAAA ATAAATTGGA AATGGTGCAG
132361   CCAGGAAAGG AGCAAAGGTC TGAGGAGGAG GAGAAGGAGG AAGAGGAGTT GTATCATTAT
132421   AAATACTTGA GGAAGAGGAG GAGAAGGAGG AGGAGGAGGA GTTGTATCAT TATAAACACT
132481   TGAGGAAGAG GAGGAGGAGA AGGAGGAGGA GGAGTTGTAT CATTATAAAC ACTTGAGGAA
132541   GAGGAGGAGG AGAAGGAGGA GGAGGAGGAG TTGTATCATT ATAAACACTT GTGACGGTCC
132601   CAGCCCCAAG ATATAGGCAT GCTAATAAAC TGAGGCTTAA CACTTTGACT ACAGAATGCT
132661   GCTTCTCCCT AACACCATCA AGGCTCCAAC TGAATAACAA TGAATTATGA ATGAAAGAGC
132721   TGTAAGGAGA GACAAAGTT  AGAATGAGAC AAGTATTGTT ATCTAGAGAT GCCAAGAAGG
132781   CAAGGAAGAT AACTAAAAAG GCACTCTGGA TTTAGAAATA GGAAGTCATT AGTGACCTTG
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132841    TAAATAATGG AGCCAGAGGA ATACCAAGGG CAGAAGCCTC ACTATAGTGT GTTGCACCTG
132901    TCAGAGGTCA GGAGGTGTAA CTGACTCTCC CACAGTGTGG CTTTGGAAGA GAGAAGTCAG
132961    CAGCTGCATG GAGATTTGGG AGAGGGAAAG CTTTTTTTTT TTTTTTTTAA TTGGAAAAGA
133021    CTGAGCTATG TGTAAATAGA ATAAGACAGG AAGAGTGTAG ACACAGGAAA GAGGGCAGAC
133081    AAAAACAAGT GCACAGTTAT CTAAGGGAAA CAATGGGATC AAGCTGCAAG TATATAAACT
133141    TGTCTTGATA GAAGAATCCT TGATCTGGTT TATTCAGTGT TTGGTCCAAA CCCACATCCC
133201    TGTTCTGCCT GTCTCTGACT TGCTCTGTGC CCCAGAAGCC CAGCTTCTAC AGATAGCATT
133261    AGCTGGGCAG CCCTGCCCTC TTGCAACAGC TGGATTTGGC CAGTGATCAG CCCAGCAGGA
133321    ATGTAGATGG CAAAGGAGAG AGAGGTTAGT GTACTTATTC CCTGCATCAC CCCCCTGCTT
133381    GGTGGGCAGC TCTTCCTCCA CAGTCCCAGC TCTGGCCTAG CTCTGGTTAC AGGTTCCCTC
133441    CCATTGCCTC TTCAGATTTA AAGGTGTGTC TGTCAGGGTA TAACTGGGAG CTAGAAATTG
133501    CACTGAAATT GAACAAAGAA TTTTATGGGA ATGGTTGTTA ACTAGTTATA AGAGGACTGA
133561    AAATGGAAAA GTGGAACAAA CGTATCAGAG ATAGTAATGA CAGAAAGCAA CTACCACCTC
133621    CAGGTTTAGG AGAACAAGGA AAAGATTCTT TGAAGAGATC CCCAGAACTG GGACCTCTGA
133681    GGAGTGTATG CTGGACCACT GATGATGATA TGTCTGTAGA TAGAGGCATG ATGAGGCTGA
133741    TTTTAGGAGC ATGGAAGATC TCCAAACTGA AGCCAACTGC TGTTACTGGA TTCAACTGCC
133801    ACTGCCAGGT TGAAGAACCC ATTCTGTGAG GATGTCAACA AACAAAGTGG GAAATCTTTT
133861    CACATCCTTC CAGCCCTCTA GTCTTCCTCC AGTGCTTTCT ATTGGTAGGG TTTGGGGAGG
133921    TGGCTAGCAA AGCGGTATTG GAAAAGATAG AAGAGACTAA ATCTTCATAA CCAGCACAGG
133981    GTGACACTGG ATCACTACTG TTGCTGATCT TGGGCTGCCT CATATCCCCT GTTCTTCCCA
134041    TTAGCCCTGT CACAACTTTG TAGATATCCC TTCATTATAT GCCCTTCATA TATTCTTTTG
134101    GTTTAACTTT TTCTGTTGGA ATCCTAATAT GGCACTCCTC CATTTTTCAG GACCAAAAGA
134161    GTATAAAGA  TTATCTTTTA CCAAAAAAAA GACAAAAAAC TGATCTAATT CCTGATTTGA
134221    TCATTACACA ATCTATACAT GTATCAAAAT ATCACATAGT ACCCCATAAA TATATACAAC
134281    TGTGTCCATT AAAAATAAAA ATTAAAGAAA AGATGGTAAA TATAGCTCTG TCAGGCAGTG
134341    GAGGTTTTAC CACGATGGCT GTTATTTCCC CCATGAAGGG GGGAGTGAGG GAGCAGCTGA
134401    AAGTAGGTGC TTATAGGGGT ATAGAGGGGC TCAAAGCTTT GAGAGAGGAG AATGTCTGAA
134461    AGAGCTGCCA AATAGCATGC AGGTCCCATG GGGGCAGAGC CTCTGCTCAT TCACCAGTGC
134521    CTCTTCAATA TCTACACTTA AGCCTAACAC AAAGTGTGTG CTTAATAAGT ATTTGCTGAG
134581    TATGTAAAGT GGAAACAGAA CCAATCTGGC AAACTTTGTA GGACTGGTGG GCAATGAAGA
134641    TCAGTCAGGT AAAATCTGTG GATATAAATT TATATTGATC AAAAAATTCA AGGTTAGGTG
134701    TTTTTCTTCA GTCATGCTCA ACGATGCTTC AGCCATGCTC AACTCTTCTG TAGCCACAGA
134761    AAAAAGTTTA CCCATAATCG AGCTGTGTCT GTGTCTGAAT AATGAAAAGA CCATGATGCA
134821    AGGGAGTTGG AGACACAGAA ACAGTGTTTG AAGTAATGGG TAATGGAAGC ATGCTACCAG
134881    GGAAAGGAAA GAAGTGGCAA TAGGAAGGAA CAGAGATCTG TGGTCCTATG TCCCCTGAGC
134941    ATATTCACAT GTTAAAGCTA ATTCAGTTTT CAATCATCAT TAAAATTTTG TTCCTAAATA
135001    TATGGCCATT ATTTTCCACA ACCACACTAA AACTTTATTA CCTCTGGCAA GTGACTATGC
135061    AAGTAACTAA GAGCAAAAAT ATCCACAACT ACCATTTGAG CTATCAATTT AGGGAAAGTC
135121    ATCTGGCTAT AATCTAAGTG ACCCTCCACT GAATGTCAGT ATCTTTGCAT ATGTGATTTA
135181    AATCTGGGCC TTCGCAACAC CATGAACTGT CTTGTCTTG  AATATCCAGA TTGAAGGAAA
135241    TAATCTGAGT AGTTACGAGT CCTGAAGCTA GAAAGATGGA AACCCCATTT GCTCATCAGA
135301    AAGCCTTAGA GCTTGGGCGC TGGCGGGTCC TGTCTCACCG GACAGAGGG  GCTCTTTCCT
135361    CCCCATCTGA TAGTCTGATA ACTAGAGAAG CCGGCCAACT TATTCTCCAA GAAGGAGCCA
135421    TCTTAGTTCC TCCTGAAATG TTCATATTTA GAAATTATTG TTTGTCAGTA ATTTAACCCC
135481    TTAATGGGCT TGCCTTGTGG TCCATACCAC TGAGTGCAGA GCTTGCCTGG AAGAATTGTG
135541    AGGGCCATTC CATCTTCCAG GCAGTAGAGT TCAGTACTTC TTTAAAATTG CTGCTGAACT
135601    CTGTATTTGA AAAGAAAGAA TCATTGGGT  GTGGTAGCTC ACACCTGTAA TCCTAGCGCT
135661    TTGGGAGGCT GAGGTGGGAG GATCATTTGA TGCCAGGAGG ACCACTTGAG ACCACCCTGG
135721    GTAACATAGC AAGACCCTGT CTTTAGAAAA AAAAAATACA ATAAAATAAA TACAATAAAA
135781    ATAAAAGCAA AAAGAAAGAG TCCATCTTAG GGACAGACTG TAACTACTCA CTGGAGCTTA
135841    CCTTTACATA GTTCAGGATC AATTATAATA AAACACTTTT GTGCAGATTC AATAGGATTA
135901    TTTTAATCCC CATCATCTCT CTGAGTTTCC AGTCAGTTTC TCTGCATGTA GACACCCTTC
135961    TCCAGCCCAC CATTGTCTCT CCTCCTATAG CTCCACCAAC AAATCAGAAC TTTTTCTAAC
136021    TGCACCTAGT GCACCTAGAG TCTACTCCAG AATGCTCATG GAGAAAGTTT CTGAAAGGTA
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136081  AAACTCTGAA TGATATTTGT AGCTAAAGGG AGACTTGCTA GAGACAATAA GCTAATAGTT
136141  GTAGACTTCA GTAGAAGAGG AATGACACTG CAATGTCAGG GTGCAGGACT TCAAGAGGGC
136201  AGAGTATGGA AACCCAATGG GAAAAATGCT CACCAGGAAC ATGAAGAGAA GGAATTACGT
136261  GTAAGGATTT CTCAATGTGT TCCCAAATTT GCCCAGCAGA GGGAGGCCTC GGGTTGATGG
136321  CAGGCTGACC ACACAATTAA AGAAGGCTGA ACCTGGGGGC TTTTAACAAC CATCGTGGGC
136381  TCTACTGTAA GCATTAGAA AAAGAAAGTT ATCCATTCAA AAATATATAT ATTTTTAAAC
136441  TTCAGAACAA AATTATGAAG AGCTATATTT ACTTTTCTAC ATTCTAATTT TTATAAATCT
136501  GAGTATATTT TGCATATATT GTTATAGTAC ATATTCAATT TTGTATTTTG CTGTTTTCAC
136561  TTAACCATTT TTACTAGATT ACTCTGTGTT CATAATAATC ACTTTTTTAA AACTTTTATT
136621  TTTATTTATT TATTTTTTTT TTGAGTCAGA GTCACACTCT GTCGCCCAGG CTGGAGTGCA
136681  GTGGCGTGAT CTTGGCTTAC TGCAACTTCC ACCTCCTGGA TTCAAGCAGT TCTCCTGCCT
136741  TAGCCTCCTG AGCAGCTGGG ATTACAGGTG TGCACCACCA AGCCCGGCTA ATTTTTGTAT
136801  TTTTAGTAAA GACGGGGTTT CACCATGTTG GTCAGGCTGG TCTCCAACTC CTGACCTCAT
136861  GATCTGCCCA CCTTGGCCTC CCAAAGTGCT GGGATAATCA CTTTTTATGC TGCATAATTC
136921  TTCAGATTTG TCAGTACGAC TGTATTTACA CTCATTGTT TTATTAGAAA GAATTCCAGA
136981  ATATTTTGGC TGCCCTAATT AATTTTACAA TTAATATGAT TTTGAAATTG GGTATTGGCT
137041  CCTTCTGAAT TGGTTTATTA AAATATATTC TAATGTAATT TATGACATTT TCATCATATT
137101  AGCATATTTA TTCTGTTAGA ATTTCATAAT TTATAAAGCT ACAAACTGTA TGTGATATAG
137161  CTTGTAACTT TATCTCATAA CTTTATGCAG TTACAAGTAG AAATAAAATG TTCCCCTCAA
137221  GATTGCTTAA AATTTATTA TAAACAAGTG TAAAAAACAA AATCACTAAA ACACTCCCTC
137281  TTTTTTCCCC CAAAATGCAT GTTTCCATTT TAACAGAACC CGTATTTAAT CAGCAGATTT
137341  CTATGGTGGC TAGATTTGTA GACTAAATAT TAAAAGTCCC AAAGCAAATG CATTTTTCTC
137401  TTAAATTTTA CTGACTTTTT TTTTTTTTCT TTTCTGAGA CGGAGTCTTG CTCTGTCGCC
137461  CAGGCTGGAA TGCAGTGGCA CAATCTCGGC TCACTGCAAC CTCCGCCTCC CGGATTCACG
137521  CCATTCTCCT GCCTCAACCT CCCGAGTAGC TGGGACCACA GGCGCCCGCC ACCACGCCCA
137581  GCTAATTTTT TGTATTTTTA GTAGAGACAG GGTTTCACCG TGTTAGCCGG GATGGTCTCG
137641  ATCTCCTGAC CTCATGATCT GCCCACCTCA GCCTCCCAAA GTGCTAGGAT CACAGGCATG
137701  AGCCACCGCG CCCCGCCTAC TGACTTTTAT CCAAAGAAAA TATAAGAGCT CTTCATCATA
137761  ACGTATGTTT CTTGCTCTTG TTATTAAATA TGACACATTT AGACTTAAAC TGATTTGAAG
137821  GTTTATGACA TTGTTTAAGT TATTACATAA TTAATTCATA AAGATAATGA CTAGTTTGAA
137881  CTACTGACAG CTCACACATC ATCAGTTGAA CAGCAGAAAG CTTATTAAGC TACTTTCTTA
137941  TGTTTCTGTC TCCCAGCTAC TAAAAGAAAC GAAACCCTTC CAGGTGTTAA GGCAAAACTT
138001  TCCTCCCCCT TTCTTCTATA AATCTGATTC CATGTTAGTG AAATTTCTAC TGATGGCTTT
138061  GGTTTCCTCT ATAGTAGAAT AGAGATCCTA TGGCAAAAGT CATGTCTGAC ATGGTAGCAA
138121  ATAGAAATGG GGAAAAGGAA GGTCTGCAAG AGCCAATGTG GGAAATGGGG AGAGGACTGA
138181  CTACAAAAAC CCAGCAGGAA TTCCAGAAGA AAACTCCTCA GGACGGGCAC ATTGGCTCAT
138241  GCCTGTAATC CCAGTACTTT GGGAGGCCGA GGTGGGCAGA TCACTTGAGT CCAGGAGTTT
138301  GAGACCAGCC TGGTCAACAT GGCGAAACCT CATCTCTACA AAAAATAAAA AAATTTGTCA
138361  GGCGTGGTGG CATGCACCTG TAGTCCCAGC TACTCAAGAG ACTTAAGTGG GAGAATCACT
138421  CGAGCCTTGG AGGTGGAGGT TGGTGAGCCG AGATCACGCC ACTGCATTCC AGCCTGGGCG
138481  ACAAAGTGAG ACGCCATCTC AATCAATCAG TCTCCTCGAA AAGCAACATT ATGGAGAGAC
138541  AGGATTCCGT CAAGGCCTGG GGCACACAGG AAAATATTAA GGCAGAAGAG AGTTTCCTCC
138601  CCACACCACA CCGTATCCCA CAGGCACTGC GGATGTGCAT ATGCAAGAGG GGTTGATCCT
138661  AAGAATTTAG AGTCACAGAG GAGGAGGCAC CAAGCAGACT GTGGAGAAAG TCATGACCAG
138721  AAAGGGACAG AATGTAAAGC TTCAGCTGAT TATCTGGCCT CAGGGATTCC AGAGGAACTG
138781  GTCCCAATGG TCTCCTGGTG ATGTAGGTTC TTAGGTTTCT TTTACAGGGG TTTTCTGGGA
138841  GATCGTTGAC CCAGTTAGCA TTCAAGCAAC TTCCACCCTG CACTTTTATT CTTTCCCCTT
138901  CACCTGCTTA GGTTTATCT GTCCAGGCAA TAATAATAAA ATTATTGAGC CCTGGACATG
138961  TACCTGTAAA GCTCCTTAAA GATGATGCCT TCTAACTCCT CATTCAACAG ATACAAAAAC
139021  ATTACAATAA AATGACTCAT GCAAGACACC CAGGTAGTTT ATAGCAGCTA ATAAAAACAG
139081  AATAACTATA AAATATGGTA AGTTTATAAA AGTTACATTG AGTATACTTT ATAAGAACTG
139141  CTTATTGAGT TTGCCTAATA ACCACACAGC ACAATAATAA TATGTATATA TTTTTAAATA
139201  TGTGTAAATA TGTGTAACAC AAACTTGTAG AAGGTATATC TGAGTACAAC CCTATTCTGT
139261  TTGGTTACCT TTTCTAGTTC ATTATGTAAG TGGCATAGCT ACCTAAGGAC TTATGCTTAT
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139321   AAATGTTACT   CAAAAAAATA   CAGAGGACAT   ATGTGGATAG   ATAATGGAAG   AGATAAGATA
139381   GGTAGGTTGA   AGGGTTGGGC   TGCCCCTCCA   CACCTGTGGG   TGTTTCTCGT   TAGGTGGAAT
139441   GAGAGACTTG   GAAAAGAAAG   AGACACAGAG   ACAAAGTATA   GAGAAAGAAA   AAAAGGGGTC
139501   CAGGGGACCG   GTGTTCAGCA   TACGGAGGAT   CCCACCGGCC   TCTGAGTTCC   CTTAGTATTT
139561   ATTGATCATT   ATTGGGTGTT   TCTCGGAGAG   GGGGATGTGG   CAGGGTCAAA   GGATAATAGT
139621   GGAGAGAAGG   TCAGCAGGTA   AACACGTGAA   CAAAGGTCTC   TGCATCATAA   ACAAGGTAAA
139681   GAATTAAGTG   CTGTGCTTTA   GATATGCATA   CACATAAACA   TCTCAATGAC   TTGAAGAGCA
139741   GTATTGCTGC   CAGCATGTCC   CACCTCCAGC   CCTAAGGCAG   TTTTCCCCTA   TCTCAGTAGA
139801   TGGAATATAC   AATCGGGTTT   TACACTGAGA   CATTCCATTG   CCCAGGGACG   AGCAGGAGAC
139861   AGATGCCTTC   CTCTTGTCTC   AACTGCAAAG   AGGCGTTCCT   TCCTCTTTTA   CTAATCCTCC
139921   TCAGCACAGA   CCCTTTACGG   GTGTCGGGCT   GGGGGACGGT   CAGGTCTTTC   CCTTCCCACG
139981   AGGCCACATT   TCAGACTATC   ACATGGGGAG   AAACCTTGGA   CAATACCTGG   CTTTCCTAGG
140041   CAGAGGTCCC   TGTGGCCTTC   CTCAGTGTTT   TGTGTCCCTG   AGTACTTGAG   ATTAGGGAGT
140101   GGAGATGACT   CTTAACGAGC   ATGCTGCCTT   CAAGCATTTC   TTTAACAAAG   CACATCTTGC
140161   ACAGCCCTTA   ATCCATTTAA   CCCTGAGTTG   ACACAGCATA   TGTCTCAGGG   AGCACAGGGT
140221   TGGGGCTAGG   GTTAGATTAA   CAGCATCTCA   AGGCAGAAGA   ATTTTTCTTA   GTACAGAACA
140281   AAATGGAGTC   TCCTATGTCT   ACTTCTTTCT   ACACAGACAC   AGTAACAATG   TGATCTCTCT
140341   CTCTTTTCCC   CACAGGAGGT   GATGGCCGGA   AGAACATGGC   AGAGGGCAAA   ACAAAACAGC
140401   ATTGGGAACA   AGCTCTGTTT   AAAAGGAGAC   TTGTGAACAG   CAAAGAGTAG   AAAGGGTTCT
140461   CTTACAACTG   AAGCCCATGG   AAGACAAATG   TGTACTGCGT   GAGTTTTAAG   GCAATAGGAG
140521   TAGTGGGACC   TAGGGCACAC   CAGAGAGCAT   ATTAACTCTC   AAACTTTTAA   AAACATTATA
140581   TCTGCTGGAC   ACAGTGGCTC   ACACCTTAAT   CCTACAACTT   TGGGAGGCCG   AGGCGGGCGG
140641   GTGTAGCTTG   AGCCCAGGAG   TTCGAGACCA   ACCTGGGCAA   CATGGCAAAA   TCCCGTCCCT
140701   ACAAAACAAA   CAAACAAAAA   ACAAAATTAG   CCAGGCACGG   TGATGCGTAC   CTGTGGTCCC
140761   AGCTACTCAG   AGGCTGAGGT   GGGAGGATCG   CTTGAGCCCC   GGGAGGTTAA   GGCTGCAGTG
140821   AGCCATGATA   ATGCCACTGC   ATCTCAGCCT   GGGCAACAGA   GGGAGAACCT   GTCTCAAAAC
140881   AAAAACAAAA   ACACACCATA   CCCAACCACA   ATGCATCTGT   CTTAAGTACC   AGTACCACAC
140941   CCCTCTACTC   ACTACTAAAT   AGGTGAGTTC   CCAATCCCTG   GTAGCAGGTT   TAAGCATGTT
141001   ATATTAAAGG   TCTTAGGCTA   GTGACTCATT   CACTCATTAA   ACAAATACTT   ATTGTGCATC
141061   TACTATAAAC   TAAGTACTGT   GCTAGGTACA   AAAGCAAATA   ATCTAAGCTC   TATAAACTTT
141121   ACTTTCTTCA   TCAACAAAAT   GGAGATGTTT   TAGGCATCTA   CTCATCATTC   TGAGCTCCAT
141181   CTTTTGTGAC   TGTAGTTGGC   AGAGCTTTTT   ATCAGTTTCT   CTAAATAGCT   CTACCAGTCC
141241   CTGGTGGATG   CTGGCATGCC   CAAAGGATCC   ATCCTGATGG   CCCTGTCTGC   TTACCTTACC
141301   TGCCTGCCTT   TGCAGCACCG   CTCTGCTCTT   CTGCAGGACT   TCCCTTATCC   TTTGGGGTCT
141361   TGCTGCTCTT   AGGCTGCTCT   GCTTGTTTTG   ATCTGCTTTG   CATCACATGT   ATGTAAAGGT
141421   CCTTTCCTTA   TTTACCCATG   ACCAAGGTAT   TATGAGATTC   TGGAATTTCC   CCAAACCACA
141481   TTGATTGCTG   GGAGAATAGA   AGAAGTGGAT   TACAAGTGGA   ACTTAGAAGG   GGAGTATTCG
141541   AGAAGACGTC   TCTGCAAATC   CATTTAGAGA   GACCTTTCTC   CAGTGGTGAC   TCAAAGATGC
141601   AGCTCCTTTC   ATCCTGTGGC   TTGGCCATCT   TCAGCACATG   GCTCCCAAGG   ATGTCCTCAG
141661   GATGGTCTCT   AATCCAAGGA   GCCTGAAGAG   AAAAAAAGGC   ATGGAGTATT   GTGAGTGGTA
141721   GGTGGTTATG   GACCAGTTAT   GGAAGAATAC   ACATCACTTT   TGCCCACCTT   CTACTAACCA
141781   GAACTCACAC   AGCCATAGAC   ACTGACAAGT   AGGACTTAAC   AAGAATCTAA   TTTTGAGTCT
141841   AGGAATACGA   CTGTAGCAAA   TATTTAACAG   CTTCAAACAC   AGGTGCATTG   CTATCACTAT
141901   GCTTGGCCCA   GGCCTGTCTC   CCTTTCCTGC   CATGTCACAG   GGGCCAGCAT   TTATGTCTAG
141961   ATTGGGTTGG   TTGGGATATT   AAGACAATAA   TGAACCAATA   CAACATCTTG   AGCATAAAAC
142021   CAACTGATAC   AATGATGTAC   AAGTCAGATG   ATTCTGATGA   TTATGAATTA   TGTCAATAAA
142081   AGAAATGTGA   TAACTAAGGT   AATTTTTGTT   TTGGCAAATT   TTTGTTTGTT   CATGACAGGA
142141   TGAAATCCTG   TCATTTGTAG   CAACATGGAT   GGAATTGCAG   GATACTACAT   TAAGTGAAAT
142201   AAGCCAGAAA   CAGAAAGTTA   AACACCACAT   GTTCTCACTT   ATATGCAGAA   GCTAGCTAAC
142261   TAAGTAAATA   AGTTTATCTC   ATTGAAGTAA   AAAGTACAAC   AGAGATTACT   AGAGGCTGGG
142321   AATGGTAGGG   GAAAGAGATG   ATAAAGAGAG   ATTCATTAAA   ATAAGTTACA   GCTAGATAAG
142381   AGCAATCAGT   TCTAGTGTTC   TATTTGTACT   ACAGAATGGC   AATAGTTAAC   AGTAATAAAT
142441   AATTTCAAAG   AGCTAGAAAA   GAGGACATTG   AATGTTTCCA   ACACAAAGAA   ATGAGAAATG
142501   CTTGAAATAA   TGGATATTCT   AATTAATTAC   CCTGATCTGA   TCACTATACA   CAGTATGTAT
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142561  AAAAATAACA CTATGGGCTG GGCGCAGTGG CTCACACCTG TAATCCCAGC ACTTTGGGAG
142621  GCCAAGGTAA GCAGATCACT TGAGGTCAGG AGTTAGAGAC CAGTCTGGCC AACATAGTGA
142681  AACTCCATCC CTACTAAAAA TACAAAAATC AGCCAGGCGT GGTGGCATGT GCCTGTAATC
142741  CCAGCTACTC AGGAGGCTGA GGCAAGAGAA TTGCTTGAAC CCAGGAGGCG GAGGTTGCAG
142801  TGAGCCGAAA TCGCGCCACT GCACTCCAGC CTGGGTAACA GAGCAAGGCT CTGTTTCAAA
142861  AATAAATAAA TACATAAATA AATATTTTTT AAAAAAAGAA CATCACTATG CACCCCATAT
142921  ATACATATAA TTATTATGTC AATTTGAAAC ATAATTTTGA AAAATGAAAA AATGAAACAC
142981  AAATATGAAT CAATCCTCTC CAAGTTGATA TACTTAAAAG GAAAAAAGTC CGAGGGCTTA
143041  AACTATTCAA TCAAAATTTT ATTAAAATGC TATAGTAATC TGGAAAGTAT TTCAGAATGA
143101  ATTGGTATAA GGTTAGACAC AAAGATCAGT GAAACAAAAT AGAGAACCCA GAAATAGATT
143161  CACACATCTA TGGACAACTG GTTTTGACAA AGGTGTCAAG GCTATTTAAT AAGTAAAAAA
143221  ATCGTCTTTT CAGTAAATGT TTCTTGAACA AGTAGACATC CGGTGTGGGG GAGAGGAGCA
143281  GGAGCCTTAC CTCAAACTTT ATGCAAAAAT TAACTCAAAA TAGACCATAG ACTTAAATGT
143341  AAAAGCTAAA ATTATAAAAC TTCTTTAAAA AATAGGAGAA AATCATCAAC ACCCTAGGAT
143401  TAGCAAAGAT TTCTTTAAAA CAAAACAACA GGTTTATAGT TTATAAAACA TAAATAACAA
143461  AATGATAAAT TTCATCAAAA GTGAAAATTT GCTTTTCAAA AAACATTATA AAATGAAAAG
143521  CAGGAGGCTG AGGCATGAGA ATCACTGGAA CCCGGGAGCT ACAGGTTGCA GTGAGCCAAG
143581  ATGGTGCCAC TGCACTCCAG CCTGGGTGAC AAAGTGAGAC TCTTCCTAAA AAATAAATAA
143641  ATAAATAAAT AAATAGAAAA GAAAAGAAA AATCACAGGC TGAGAGAAAA TATTTATAAT
143701  ACATGTATCT GACAAAGGAC TCGCACCTGG AAAATATAAG GAACCTTATA ACTTAGTAAG
143761  ATGACAAGCC AAAACAAAGA GTAAAGTTT TCAACAGACA TTTCACAAAA GAAAACATAC
143821  AAATGGCCAG TATGCACATG AAAAGATTTT AAACATCATT AGTTACTAGG GAAATGCAAG
143881  TCAAAACCAC AATGAGATAC TTCACATTCA ACAGAATAGC TAATGTTAAA AGGACTGACA
143941  ATCCCCAGGG TGAGCAAGGG TGTGGAGGAA ACTACTCTCA TATATTGTGA ATGTAAGAGG
144001  CATTTTATGA TATAACTGAA TTCAGTTTTA TGTATAACTG AATTACGGAT ATGAGAATCT
144061  CAAATGAGGA CGAATGGTTT TTACGCACAA AACATGAGAC ACAAATCTGT AAGAAATATA
144121  AAGTCGTGAC CACGTCCTTT CAGAACTTTA ACCTGTTTGC TGAAGTACGT CAGTAACAAT
144181  GGCAGGGAAA GGGTATCTTA AATTTCACCA CAGCCTCAAA GAGGCCATTT CGTGGATCCG
144241  CTGAGGCTTG GAGTCGGCCT TCTGACCACG AGTCCTGCGG CTATGAAAGA GGAAGCCGCG
144301  GTTCAGGGCG TCCTCGCGAG TCGCGCAGCC CGCCCTGCTC CAGCTGGGGA CACAGGTGGT
144361  CACGGCGCTT TCCAGCTGCA GATCCAGGCG GCAGCCCAAG ATTTGGTCCA GCCGCCAAGG
144421  GGTGGCTCGA GTGACTGACG GGCCTTGAAC GCTCCCAGGA CCCACATCTG GAGAGGGAGG
144481  TGGGGGTGGG GTGCTGAAGT CATTCTTGGG GCCCCTGGGG GCGGGCATGG ACCTGGGTAA
144541  GGCCAGAGAA ATTGACACCT CGTGACATCC CTGGAAGAGA AGTACGTTCA GTGTCACTCC
144601  AGAGCTGAAA GATACCGCCT TCTGGCTGGT CCCTCCTCAC CTACATACTT TTCTAATTTG
144661  TCTGGAGCAG GCCGGGCATC TGTATTATCT GGTTATTTAA ATATCTGGTT ATTTAAAAGC
144721  TCTCCATTAA ATTCACATAC ACGAAAATAA AAATTAAAAA AAATTTTAAA AAAAAGAAAC
144781  AAAAGCTCTC TAATGACCAA GTCCTACACG ATAGTGAATA AATTTTTTTG TGTGGTCCCT
144841  AAAATTGAGT TCATGCCTTT TCTGAAGTAA TAGACGCCCA GAGAAGGGAT CGACTTACCC
144901  ATCATGCCAC AGAGATTAAT TGGCCCCAGA ATTCTTTAGC AGACCGTGTA TATGAACGTC
144961  CTTTGCAATC ATATAAATTA ACTGGGAAAA CCTCATTTAG TATGTTACAT GCCTAGCGTT
145121  TTGTGCCTGA ACACCTTACA AGAACCAGGG ACTATTGCCC CAATATTATA TTTCAGGAAA
145181  GGAAGGCCCA GACAAATGGT GTCACTGGTC CACTTTCACC CAGTTGGTAA ATGAAACCAG
145241  AAATTATAGC TGTACCACAG AAAGGTGAAA ACGTTTCTTT TATAATTTCA CATACAATCT
145301  TTAATGGACC CAGTGTCCAA CACATTAAAG CAAGTGCTCA GGAGTGACAT CAAGATGTAA
145361  AAAATAGTCC TGTCCTCAGG GAGTTTAGGT CTTGGAGAAA AGAGACCCAA GGAGACACAA
145421  GACAAAGGGG AAAGAGAAGG AGCGCTGAAG ACTGAGGACC CTGCCTGTGG ACTGAAGTGA
145481  GGATGGGGAC ACCCGATGCC CGGAATATGA CAGTTTGGAG GGGCCTGAAG GACTCTTCTA
145541  TTCTCTATCA GAAAAACAGA ATTACTCTCC TAACCAGAAA AGGTATTTCA ATTTATATTT
145601  TCCATCACAG CACTTTTCTG GTGATAATTT AATGTGTTTT AAAAAATGTA TCACAGTGAT
145661  GGCCTGGTGT GAAATAAATA ATAAAATTTT AAGAATTAAA AAATATAAAA ATCTTTTATA
145721  TAGACATTAG GAGTTACAAG GATAACTGTG AATTATAATT AGTAATTAAA TTGAAATACT
145781  GATTATTTTC ATTTTTATTT AATTATTTAA TAAAACCTAT TTAACATTTA ATATTTATCA
145841  GTAATTAAAT CTAATTGTTA ATATTTATTA TTATAAATTA TTTTAGAATT AAAAATAAGT
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145901  GTAGAAGCGA  GGCATGGTGG  CTCAAGCCTG  TAATCCCAAC  ACTTTGGGAG  GCTAAGGTGG
145961  GAGGATTGCT  TGAGCCCAGT  AGTTCAAGAC  CAGCCTGGGC  AACATGGAGA  AACCCTGTCT
146021  CAATACAAAA  AAATGAGCCA  TGTGTGGTGG  TGCGTGCCTG  TATTCCCAGC  CATTCTGGAG
146081  GCTGAGGTGG  GAGGATGACT  TGAGCCTAGG  CAGTCAAGGC  TGCAGTGAGC  CCTGATCTTG
146141  CCACTGCACT  CCAGTCTGGG  CAACAGAGCA  AGACCCTGTG  TCAATATACA  TATGGACAAA
146201  CTTAAAATTT  AAAATGAAAG  CATACTACTG  ATACAGAATT  GAGTAGAGAT  GCAAAGCTAG
146261  TCCTATAACC  AGAACAATAA  AGATAAAAAG  GAGAGTGGAA  GAAGGTATGT  CATGAATTTC
146321  ATGATAAATG  GCAATTGCAA  ATATCCTGTA  GCAGAACAAA  ACAACAAAAT  TGTAGATAAA
146381  ACATATCCAA  CCCTTTGGAA  GGCCAAGGAG  GGAGGATTGT  TTGAGCCCAG  AAGTTGGAGA
146441  CCAGCCTGGG  CAACATAGTG  AGACCCTGTA  TCTAAAAAGG  AAGAAAGAAA  AAAAAAAAAA
146501  AGGATGATAA  AGTAGACAAT  ATTGAAAGCC  ATTTTCTGCA  AATACATAGT  GAATTTGATC
146561  AGTAATTTTC  TTCCAACAGT  GCAAAAATGA  ATAGATATTA  GTTGCCTGAA  ATAAAAATCA
146621  AATATCCAAC  AAAAAATATT  GACTATCTAA  TAGTATCTAA  GCTAGTAAAT  TTGGCCAGTT
146681  ATAAAATGTC  TTAAATTTTT  ATTTAAAAAA  AGAAAACCAT  ATTTATAAGA  AGAGGTGATA
146741  AAGAGAAATT  ATTTCAGTTA  TGAAGATTTT  GTTAGAAAAC  TATGAGAAAA  AAACTATTTT
146801  TTGTTTTCAA  AAAGTGAAAG  ATTAAGTTAC  CAAACAGTTG  CTAAAGAATA  CCAGATGGCT
146861  GAGCGTGGTG  ACTTATGCCT  GTAATCCCAG  TACTTTGGAA  GGCCAAGGCA  GGAGGATCAT
146921  TTTAGGCCTG  GAGTTCGAGA  CCAGCCTGGG  CACTGTAGCA  AGACCCGTCT  CTATTAAAAA
146981  AAAAAAAAAA  AAAAAAAAAG  AATACCAGAC  CTTGCTAACA  ATAGCAAAGA  TCAATTAATT
147041  CAAAATTTGA  AAAACTGTAA  TTTATTAGC   TTTAGAGTAC  TCTCGTGATA  TGAGATTGCC
147101  AAATTAATAC  TTTGGGTGCA  TTTCTTTTCT  CAAAGGACTT  GCAAATTTAC  AAAGAAGTGT
147161  TGAAGAAAAG  CCACACATTG  GCAGGTAATG  TTTGCAAAAG  ACAGATCTGA  TGAAGAACAA
147221  TATTTTTAGA  ATATACAAAG  AATACTTAAA  ACTCAACAGT  AAGAAAATAA  CCTGATTTAA
147281  AGCAGGCCAA  TGACCTGAAC  ATCTGTTCAC  CAAAGAAGAT  ACACAGATGC  AAGTATGCAT
147341  ATGAAAGAT   GCTTGACATC  ATGTCATTAG  GGAACTGCAA  ATTAAAACAA  GTAGATACCA
147401  CTGCATACCT  AGTAGAATGA  CCAAAATTTA  GAACACTGTC  AGCACCAAAG  GTTGCAAAGA
147461  TATGTAGCAA  TAGTAACTTG  TTCATTACTG  GTGAGAATGC  AAAATGTGCA  ATCACTTTGG
147521  AAGACAGTTT  GGTGGTTTCT  TACAAAAGTA  ACCATACTTT  TACCATAAGA  TTCACCAATC
147581  ACACTCCTTA  GTATTTATCC  AAAGGAATTG  AAAACTTATC  TCCACACAAA  AACCTGCACA
147641  TAGATGTTTA  TAGCAGCTTT  ATTCATAATT  TATCCAAAAC  TTGGAAACAA  GATGTCTTTC
147701  AGTAGGTAAG  TGGATAACTG  TGGTACTTCT  GAATAATGGA  ATGTTATTTA  GAGTTAAAAA
147761  GAAATGCATT  CACTTTGGGA  GGCCGAAGTG  GGTGGATTGC  TTGAGGCCAG  GAGTTTGAGA
147821  CCAGCCTGGT  CAACATGGGA  AAACCCCAAT  TAGCCGGGCA  TAGTGGCGTG  AGCCTGTAAT
147881  CCCAGCTACT  CGGGAGGCTG  AGATATGAGA  ATCGTTTGAA  CCTGGGAGAT  GGAGGTTGCA
147941  GTGAGCCAGT  GCCACTGCAC  TTCAGCCTGG  GCAACAGAGC  AAGACTCCTC  TGTCTCAAAA
148001  AAAAAAAAAA  AAAAAAAAAA  AAAAAAGAA   AGAAAGAAA   AAAGAAAAG   AAAAAGAAAA
148061  GAAACGATCA  AGCCATGAAA  ACACATGAAG  GAAACTTAAA  TGTATGTTAC  TAAAAAGCCA
148121  ACCTGAAAAG  ACTGCATACT  ATATGACTCC  AACTGATGCA  GGGCAAGCAA  GCCAAAAATT
148181  AGGGCTTAGC  CCGGGAAGAA  TTCAAGGGTG  AAGTGGTGGT  GTTAGCAACT  TTTACTGAAG
148241  CAGCAGTGTA  CAACAGCAGA  ACAGGTACTG  CTCCTTGCTG  AGCAGGGCTA  ACCCATAAGT
148301  AATGTGCCCA  GAGTAGCAGC  TCAGGGCAG   TTCTGCAGTA  ATATACCTGC  TTTTAGTTAA
148361  GTGCATGTTA  AGGGGGATTA  TGCAGAAATT  TCTAGAAAAA  GAGTGGTAAC  TTCGGAGTAG
148421  GTACAGAGGA  AAGAAGTCGA  TAATGTCCTG  TTGTTGCCAT  GGCAACGAAA  AACTGACATG
148481  GCGCTGGTGG  GCGTGTCTTA  TGGAGAGGTG  CTTTAACCTC  GTCCCTGTTT  CGGCTAGTCT
148541  TCAATCTGGT  CCGGAGTAAA  GTCCCTGCCT  CCGGAGTTCA  CTCCTGCTTC  CTGCTTCACA
148601  ACTGTATGAC  ACTCTAGAAA  AGACAGTAAC  TATGGACACA  GTCAAAGAT   TAGTTGATAG
148661  AAATTGGGTG  ACAGGAAGTG  TTGAAAAGGC  AGAACACAGG  ATTTTTAGGG  CAGTGAAACT
148721  TCTGTGATAC  TATAATGGTG  AATACATGAC  ATTATACATT  TGTCAAAACC  CATAGAAAGC
148781  ACAACACCAA  GAATAAACCC  TAATGTAAAT  TACAGACTTT  CGTTGATAAT  GACGTGTCAA
148841  TGTAAGTTCA  ATTGTAATAA  ATGTACTACT  GTGGTGCTGG  ATGTCTATGG  TGGGGGGACA
148901  TTTTTGCTTC  AATAGTTACA  GTTGAAGTAA  ATGTTTGTGT  TTCCCACAAT  GCATATGTAG
148961  AAACTCTCAC  ATTCAATGTG  ATGGTCTTTG  GAGGTGGGCT  CTTTGGGTGA  TAGTTAGGTT
149021  TAGTTGAGAT  CCTAGCAGAT  CGAGTCTTCA  TGATGGGCAT  GATGGGACTG  GTCCCTTATA
149081  AGAAAGACC   AGAAAGCTAG  CTCTCTCTTT  GCCATGTGAA  GACATAGCAG  GAAGGTAGCC
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149141    ATCTGCAAGC TAGGAAAGGG CCTTCACAAA GAATCAACTC AGACCTCAGA ACAGTGAGAG
149201    ATAAATTGTC GTTGTTTAAG TCACTCAGGC TGTGGTATTT TGTTTCAGCA GCCCAACCTA
149261    AGACTGTTAA TTGGATTAGA AAATTTCCTTT TGGGGATGGT GTGTGGCGGG GGGTGCGGGG
149321    AGTACCTTTG TTAAGCTTTT ATATCAATGA GTTTGTAGGC TTTTCTTTTT TGGTCATTGA
149381    CTAGGACAGT TTAAATAGTA TGAGTGTGAA GGAGATTGTT GGTCATCTAT TCGATGTCCC
149441    TTCTCTGTTT TTTAATATGA GAACTCCTGA TTTTCAGCCA ACTACCCTGG AAAAAAAGCT
149501    AATCTTTCTG ACTTCTTAAG TGTGGCCATG TACTAAATTC TGGCTAATGC AAGGCAAGCC
149561    AAAGGTTTTA TGATAGGTTT TAGGACACTA GAGTAAAAGA GAGCTGTTGC ACACATGCTC
149621    TTCACCCTAC TTTTGTGTCC TTTTTTCCAT CCTACAACTT GGGTTGTGAG TATGATGGCT
149681    GGAACTTTAG TGGCTCTCTT GGATCCCAGG GGTAATTGAG GGGTGGCTGG AAGGAATCTG
149741    TGATTTTCTG GAGTTTCCAT ACACAAACAA GACCTGGATT TTCTGGGCTT CCCAGACTTC
149801    CACATCTAGA CTTGCTTTAA ATGGGAGAGA AATAAACTTG TTTCAGCCAC TGTCATTTTG
149861    GGCTATTTTA TAGAACTTAA TCTAATCTTC AAGGGTACAT GAATTGCTTT TCCTTAAAAA
149921    AAAAATCAGC CATAAAATCA TCTTCTTTTT TCTTTTGTTC CCCACATTAT TTAGTTGGAG
149981    CTCTGTAACT TTTTTTTTTT TTTTTTTTGA GACAAGGTCT TGCTCTGTCA CTTAGGCTGG
150041    AATTCAGTGG CATGACCATG GCTCACTGCA GCCTTGCCCT CCTAGGCTCA AGCAATCCTC
150101    GTCTCAGCCT CCTGAGTAGC TGAAACTAAG GCACATGCCA CCATGCCCAG CTAATTTCTT
150161    TTCTTTTAGA GATGGGAGCC TTGCCCAGGC TAGTCTCAAA CTCCTAGCCT CAAGTGATCC
150221    TCCCATCTCA GCCTCCCAAA GTGACAGGAT TACAGGTGTG AGCCACCATG CCTGGCTGCT
150281    CTGTAAGTGT CTGAATTTCA TTTTGTATTT ATCAGTCTGT TTAGATTTTC TTTCCCTTCT
150341    TGGGTCAGTT AGGCCATTGG TTTCTTTTTA AAGGTTTTCA AATTTATTTG CATCTAATTC
150401    TTCAAATTAC TCTCAAAATT ATTCCAGTAT ATATTCTTTT GTTCCTATTT TCTTCTGTAT
150461    TCTTTATTAA AATAGCTAAT GATTTATCTA GCAGGACTTA TATTCTTTCC ATAACTTTCC
150521    TGCACCCCAA TTAATCTCCA ATTTTATATT TCTTCTGGCC TTCCTTATAG TTTCCACAGG
150581    TTTATTTTAT TCATTTTTTA AAACTTTTAT TTAATTGTTT ATTTTATTAT CATTCTTTCT
150641    TATTCAGCAA TCTAAGTGCT TAGGGATATA GAATTTCCTC TAAGCAGCAT ATGCTAGGCT
150701    TTAACAATGT TAGGGAGGCC TCCCCTTTCT GGGGAAGACC ACACTTACAT TAACACAGGA
150761    CTGTGGGATG CCAAGAGGTA GAGAAGAGCT TATGAATATC CAGATTACAT CTTCACTGAT
150821    CCTGCACAAA GGTGGGGGTTC CTCGGTTACC CACTGGGTCC TATTACCCAA GTCTGGGTCA
150881    GCATACCGAG ACTACGGGTA TATAGAACAA GTGCAACTGG CGATAATCCT TCTGTTGGGG
150941    AGAAAAATCT TTTTTTTCTA TTCATCTTAG GTTCTCCATC TGTGGCCCTA TCAAGTAGAC
151001    TAACAAAAGA CAGATTGACA AGACAGAAAC AAAGCATGTG CATTGTACAA ACACAGGGGA
151061    GTACTGAGAT GAATACTCAA AAGAGGATTT AGAACTTGGG CTTATATAGC ATTTTAAGAA
151121    AAGAATACAT TTTTTAAGTG ACAAGGAAGA CGAAAAGGAC TTTGAGTTTC TAGTGCAGTA
151181    AATTGTGGGA AGGCAACTTT TTCTTTCCCT TTTTTTTTTT TTTTTTTTTA AAAAAAAGAC
151241    TTCTCTGGTG CTATGTCCAG GCTGATAAGA GTCTAAAGTC TCTGGTGACT AACTTTTGTT
151301    CTTCCCCGAG TAAGAAGACA CCTTCACAAT TTCATATCCT GCTTTTAGGC AAACAGGGAG
151361    AGGGCAGAGG TGTTTGTTTG TTTTTAATCT ATTTTTTTTC TCAATTGTCT TCAACTCAAA
151421    ATACTTCTTA TGCCAAAGAT GGCATATTCT GCTACCCTTC ACTTACTACT TACAACCCAG
151481    CCTCTATCAT CATAATTAGA ACTTCTGACC CTGGGGAACA TGGGCAATAG TTTGAACTCT
151541    TTTATATCTC CCTTAGGCAG AGATGGAGGC CCAGCCATGC CTCTGACATC TAGACACAAC
151601    TGTTGCTTCA TTTCTCCTAT TCTCAGAGGT GATGTTGTAG GACTTCAACA AATATCAGTA
151661    AACATTAATT TTTTTTTTCC TTGAGGCACA GCATGATCTT GGCTTACTGC AGCTGCTGCA
151721    GGCTCAAGCA ATTCTCCTGC CTTGGCCTCA CGAGTAGCTG GGTTACAGGC CCCTACCACC
151781    ATGCCCGGCT AATTTTTGTA TTTTTAGTAG AGACAGGGTT TCACCATGTT GGCCAGGCTG
151841    GTGTTGAACT CCTGACCTCA AGTGATCCAC CTGCCTCAGC CTCACATAGT TCTGGGATTA
151901    CAGGCGTGAG CCACCATGCC TGGCCATCAA TTTTTATGTC AACTCTAAAT TATAACATTT
151961    AGCAATTTTG TGACTTTTTA TGGTCATCAT TAATGTTGTT TATGTTTTAG TTGTAGTCCT
152021    GTCATTACTC ACTCGGGTAT GGTAATTTGG TCTTTTTCAA AATGAAGTTA AGGTCTATTT
152081    GCTCTTCTCT GAATCATAAT AAGAACTGCC AACAGCCATT TCAGCAATAA CTATTTACTG
152141    AGATTTTAAA ATATTTCAAG GTAATTGGTC CTAGCAGACT GGAAAATACC AAATTCTTTT
152201    CCAGAACTGA ATCCCCCATC AAAGTTCAAT TTTACTCATA ATTCCCTTTT CATTTGAAGC
152261    ATCTCATTGT AAGCCAGTCT TAACCCTTCT CTCACACTTT GCTTGGCTGT TTCTCAGGTA
152321    GAACTCAGTA AGTCTGGTAG CCTCCAGGAC TGCCGCTTAG ATTATTAAAC AACATGTCAG
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152381   TGGTTGGAAG AGTCAATGTT ATTTTGATTT TTCTGTTTTG TTTTGTTTTA AATGCAGTTG
152441   GCGGATAATT GCAGCTTTCT TTCATTCCCT ACATGAGTTC AAATGGCAGC AAACAAACTA
152501   GGAGAACGCA GACCTTCTGA CTTGTGGGTA CCCCTACTCA TCACCTGAAG ACCCTTGGAA
152561   ATCAAAGCCC TGACCCATTA AAGACGGATG GAGACAGCAA CATACGATCA TCACTATTAT
152621   CTTGCTTTGC CCCAGTCCAG GTTAACCATC TGTGGTATTT TTAGTTGCTA AGTCCATATA
152681   TTCAACATAA ATCAATTATA TATCCACTAA AATCTCAGCA CTAGTCTAAC TACTAAGGAA
152741   ATGACAGCGA AGAAAACAGA CCAAACGTCT GCCCTTATGG GATTTATATT ATTTTCTCTG
152801   TGCTGGTTAA ACCAAGGAGC TTCTGCTCTT TTCCTTAGTC ACCTGGGGGA GGCAGAAACA
152861   AAGGAGAATA TTGATAAACC TGGAAATAGG GCCGGAGAGT ATCAGAGAAG GAAGCCTTCG
152921   GGAAAGTAAA GATGTGGCAG CCAGTATTCC CGTTATAAAA GGATACAACT CCGGCCTCAT
152981   AGTCCAGAAA AATTCCCACA AGCAGGGGCT GCTCATGCAG ATGAAGGGAA GTTGGGGGAG
153041   AAGTAAGTGC TACATAGCCT TTCTTTTTGC ACAGCCTGAG GGTCCAGAAT CCAGACTGAG
153101   GCTCTTGCTT CATGCCAGTG CCCCTCTGCA CATTTTCCAT ACAAACTCCT AAATCCCATC
153161   CGGTTCCTTC GCCAACATCC ACTTCAAAGT AACGTCTTCC TGAGGTGAAG CCTTCACAAC
153221   CCAAGACACA GGGGAAGGCA GTAAATCTCC TGGAAGATGT GTCCTGATTC TCCTGGGTGT
153281   ATCCACGAGT CACTTGTCTC CGATCCTCAG AGAGAATTAG TTCGTGATGA GCTGTATCTG
153341   GATCCAGAGT CACACTAACT GCAAACAAA ACAAAACAAA CAAAAATAAT TTTGTTGCTG
153401   TGAAGAACAC AGGTTATTTT ATTTTATTTT ATTTGAGAT GGAGTGTTGC TGTCACCCAG
153461   GCTGGAGTGC ACTGGCACTA TCTCAACTCA CTGCAACCTC CACCTCCTGG ATTCAGGCAA
153521   TTCTCCTGCC TCAGCCTCCG GAGTAACTGC GACTACAGGT GCGCACCACC ACAAGTGGCT
153581   AATTTTTTTA AATTTTCTGT AGAGATGGGG TTTCGCCATG TTGGCCAGGC TGGTCTCAAA
153641   CTCCTGACCT GAAGTGTTCC ACCCACCTCG GCCTCCCAAA GTGCTGGATT ACACAGGTGT
153701   GAGCCACCAT GCCCAGCCAC AAGTTATTTT CAATAAAACC AGCCTGTGTT CAAACCCAAC
153761   TATTGTTTCT TATAAACTGG GTGAGCTTAG GCAAATCATT TAACTTTCTG AGCCTCAGTT
153821   TGTTAACTAT AAAGTGGAAA TTACCGTATT TGTTGCAGAG AATGGTGGGT AGGATTGAAT
153881   AAGCTTATGT TTGCTTAATG CTTGGTAAAA TTCCTGGTAC ATGGTAACCA CCTAATAAGT
153941   GGTAGTTGTT GGGGTGATCA GGCCCAACAC CAGGCCGTGG GGGCTACAAA GTCCGGCGGG
154001   GTCAAAGGAA TGAGAAAAGA CAAGTTAAGA GTGCATAAAG TGGGTCCAGG GTGCCAGCAC
154061   TAGATTGGAG GCTGCAAAGG CCCTAAGCTC TGGGAGCCCA CACTATTTAT TGGTGATCAA
154121   ACAAAGAAGC AGGTGGTGAG GACGTGAGGG TAAACAGGTG AGGGCATGAG GACATGGGGG
154181   TAGAAAGGTA GTGGTGCATT AAGCGTAGCT GTGACAGTTT AGCATTTTCT TTGACACATG
154241   TAGAATATAC TCTGCTGCTT GAGATAGTAG AGGACACGTT TATGAGTGAA AAGCAAGGAA
154301   CCAACAAGTC TGTGCACTTT CCAGAGGCTA TGAGGGGTTT TATGCCCTGA GCCCTGGGTT
154361   CCATCCAAGC CACAAGGGGT TTTATGCCCT AGGCTTAGAT TTGTGGTGCG GCAGGGCAGC
154421   CTTCCACCAT TTGGCACAGA GCTTGGTGTT CCAAAGGCCA CGAGGGGTTT TGGACCCTGG
154481   ACCCCGGACA TCTTCCAAGA CTCTTTTACA TTATGACAGA CAAGCCAGTC CTGCTTCAGC
154541   TCTTCTAACA ACATGTAGTA ATAATGATAT CATCAACATC ATCTTCGTCT TAATTATTCA
154601   AGGATGCCAA GGTACAGAAC TAACCTGTTA ATATGGTTAC CATCCTGTCC AAAGTTCTTC
154661   TCCCATGCAG GACTTCCAGG AATCATGAGA CAGTTGAGCA GAAAGATACC TTTTCCCTTC
154721   TCTACTGAAT AACCACCAAC ATTGAGAATC AGAGAGGGAA AATGACTCAG CTAATGTCTT
154781   AGCTTGTTAT TGGAAGACCC AGGTCTCATG ACACATGCCT AGTCCCATGA CTTTTAATTG
154841   TAAGCTCTTC TCTTTCCCCT CAGATAATGT TCCATAAGCA TTAGTATGAG ATAATAATAC
154901   ACTGAGGACC AATATACATG AAAAATATCA GACTAGAATC AAACAAGACA GAAAAAAGAT
154961   CTGATAACCT AAAGTGAGAT ACTGAACAGT ATGCAGTTTT AAAATAAAA AATGGTAATA
155021   GGATGTTCTA ACAAGAGAGT TAAGAAACCA CTGTGCTACT GAGTTAAATG TTGATCAGTT
155081   GGTCTGTGAC AATTAAGGAA TTCAAGTATT CAGAAACACT TCCTGTGCTG GATGCTCTCT
155141   GTTTGTTCTT CCAAATAATC CCTCACTTTT CCCTGTCTTG CTCTGTGCCC AGGAAGGCTG
155201   ACATGGACAG ATTAACCAGG CTTTCCGCCC TCTGGCTTGG TTCAGCCAAT GGGAAGCACC
155261   AGAGGAGACC ATAGGGCACA AAGAAGCAGC CTTGGGAGTA TTCAGTACCC CAGTCCCACG
155321   CTATGATTTG GAGGGTCTGC ATTCCTCTGC CTCTGGGCAC ACTCTAGTAT AGTTACAGCT
155381   CCCTACACCT GCCACTTGAG GCCCAGAGGA GGTGATGGCT CTCTAACTGT TCCTAGTTCT
155441   GGGTGCTTCC TGTTCCTTGT GGATTCCCA ACTCCTCACC TTTGTAAATA CCCTCCTTTT
155501   TCAAACTCTA TTCAGTTAGC TTTTATCAGC CTGACTCACA GAAGTTTGGG GTTTCAATTC
155561   ATATTACCTG AATGACCCAG GAAAACCCAT GTTGAGAAAT TAAAATGTTT ACGGGGTGGT
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155621    AATACCACTT AAGAGAAAAA ATATCAATTG GATTTTTAAA ATTCCACCTA TCTATTGGTG
155681    TGACACATCA ACAAAAACAT ATAGAAAGAT TGGAAGCTAA AAGATAGATA ATATAGTCAT
155741    ATACTGTTAT AGTATTATAT CAAAAGATAT TAAGTCAGAG CATTATTAAG AATGGAAGAA
155801    GGGCCAGGTG TGGTGGCTCA TGCCTGTAAT CCCAGCACTT TGGGAGGCCA AGGCAGGCGG
155861    ATCACTTGAA GCCAGGAGTT CAAGACCAGC CTGCCCAACA TGGCAAAACC CTGGCTCTAC
155921    CAAAAATACA ACAATTAGCT GGGCATTGTG GCACATGCCT GTAATCCCAG CTACTTGGGA
155981    GGCTGAAGCA CAAGAATCAC TTGAACCGGG GAGGCAGAGG TTGCAGTGAG CTGAGATTTC
156041    GCCACTACAC TACAGCCTGG GTGACAGAGA GAGATTCTGT CTCAAAAAAA AAAAAAAGA
156101    AAGAATGAAA GGAGTCACCT AAAAAGATA ACACAATTTT AAACATAAAT GTACTACATT
156161    ATTAGTGAAT TCATGTTTAG AATTGTGTTA ATATACAAAG CAAAAATTGT AGAATTATAG
156221    GAGAAATGGA CAAATCTACA ATCATCATGG GATGTTTTAA CATTCTTCTT TCCATAATTG
156281    ATAGATCAGG CAGACCAAAA GAAAGAAATA AGGGAAGATA CGGAAGGTCT GAACAATCTA
156341    AGAAGCGCAA TCTCATAGTC AATACATAAA GCTCAGCAAT TGTTTAATAA TAGTAAGCAG
156401    AGAATATGCA GTTTTCTCAG GTATAGATGG AACATGCACT AACTGAGTAA ATACTAGGCA
156461    GAAAACAGTC TGAACAAGTT TCAATAAATC TGTATTACAC AGATCATTTT CTCTAGCCTC
156521    AATATAAGAT TATAAACCAA TAATAAAAG ATGACTAAAA AGATTCTAAA TATTAGGAAA
156581    TGTAAACTAC TAATAAGTCA TTAGAAGATG TATAGAATGG AACAATAATA AAATGTTATT
156641    TATAAAAATA TACAATGAAG CTAAAGCAGA ATTTTAAGGA AAATTTGTAG GCTTTAAATG
156701    CTTATCTTAG AAAAATTAAA AAGCTGAACA TTAATGAGCC AAGCATCTAA TTTAAATTTT
156761    AAAAAGAACA TAGAAAGCCA AATATAATTT TTTAAAAAGA AAAAATAGAT ATTAAACAAT
156821    ATAACAGTGA AGTTAAAGAA AACAAGAATG CAATAAAGAG GAAAAACAAA CAAAAAAAAA
156881    AGTAGCTTCT TTTAAAAGAA ATTTAATAAA ATAGACATAC CTCCAATGAG ATTTATCAAA
156941    GTAAGACAGA AGGCACAAAT GGAATGAATA CAGAAACTTT TTAAATATTA CAGAACTTTA
157001    TAATAAATCT TATGCTACTA ATAAAATTGA AAGTACTGAT AAAATTATTA CTTCCTAGAA
157061    AAAATATTTC TGAGTAAAAC TCACTCAAAA AACAAATAAA GCATGGGCAG ACCTAACATT
157121    AAAGAAATGA AATCACTACT TTAAATTTTA CCGACAGATA ATAAAACGTG CATCTTTATC
157181    AAGCAAAAAT GGAACTTGTC AGTTTTATAG GAAATTTAGA AGTCAAGGCA TGAGTAATGC
157241    CAATCTCATA CCAAATCCTA CAAAGAATAG AAAATTATGG CTCCCGCTTA TAGACATAGA
157301    TATAGAACTC CTGCACAAAA TAATATAAAT AACAAACCAA ATTTTATATT TGCAACTATA
157361    CATATTATAT GTGTATGTAT TATATATGTT AACATATACA TATATAATAT GTATAGCATA
157421    TGTTCTACAT ATTATATATG TATAGTGTAT GTATTTTACA ATATATAAAT GAAAACCCAA
157481    TCTTTAATAT ATTCATCTAG ATTGTCATAT ATGACATATA TAATACATTA CATCAAAAAT
157541    GTGTACAATA ATCAGGCCAG GCACAGTGAC TCATGCCTGT AATCCCAGCA CGTTGGGAGG
157601    CTGAGGCGGG TCAATCACTT GAGTCCAAGA GTTTGAGACC AGCCTGGTCA ATATGGCCAA
157661    ATTCCATCTC TACAAAAAAT ATGAAAAATT ATCCAGGCAT TGTGGTGCAC ACCAATAGTC
157721    CCAGCTACTC GGGAAGCTGA GGTGAGAGGA TCACTTAAGC CTGGGAGGTG GAGATTGCAG
157781    TGAGTCGAGA TTGCGCCAGT GCACTCCAGC CTGGGTGGCA AAGGGAGACC CTGTCTCAAA
157841    AAAAAATTAA AAAATTAGCC AGGTATGGTG GCCTGTTCCT GTAGTCCCAG CAACTGGGGA
157901    GGCTGAGGTG AGAAGATCAC TTTAGCTCAG GTGGTGGAGC CATGATCGCA CCACTGTACC
157961    ACTCGGCTTG GGCAACAGAG TGAGAGCCTG TCTCGAAAAA ACAAATATAT ACACACAGTA
158021    ATCAATATAT ATATTATATG TACCAATCAA TGCTTCACTT TTATATATAA TATAGATTAC
158081    ATCTTATTAG ATATATAGTA TTCCTTCTCC ATAGATAGAT AGATACAGAT ATAGACATAG
158141    TATCCTCTAT CCATATTAGA GAGAGGATAC TATATATATC TATAGCATAT AGAGATGCTG
158201    TCTCAAAAAA ATTTAAACAT CAGCCAGATG TGGTGGCCCA TGCCTGTAGT CCCAGCTACT
158261    GGGGAGGCTG AAATGAGAGG ATTGCCATTG ATCCTCTCAT TGGTTGAGCC ATAATCGCAC
158321    TACTGCACCA CTCAGCCTGG GAGACAGAGG GAGACCTGAG GTGGAAGGAT ATAGATATAG
158381    ATATATAAAT AAATATGTAT AGAGAGAATA TAATATATGT GTGTATGTGT ATATATATAT
158441    ATTATGAAGA CACTGGGAGA GAATACTATA TATATATGTG TGTGTGTATA TATATATTAT
158501    GAAGACACTG GTGGGATGGT TTCATTACCA ATTGGACCAA GAGTCCAGGT ATGGAGCCAA
158561    CATGCAATGT TGTTGTTGAC TGAGCTGGCA GAGCACTGGT CATAGTTACG GGAAAAGAAG
158621    GTCTCCAATG AGACATACTT AACAAATAT ATGAACTTGC CATATACGTG GAGAGTTCTG
158681    GTGTGTATAT AGCCTTCTCT CACCAACCTA GCAATTGTCT TCATCATCAT TATAATGCTA
158741    TCAGAGCAAA GATGACAGCT AAATTTTTTT GTCCCTTTCT TCTTCTTTCT CTTCCTTCCC
158801    CTCCCCCACC TCTTTCTCTT CCTCCTCCTC CTTCATCTCT CTTCTTTTTT TTTTGAGAT
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158861   GGAGTCTTAC TCTGTCGCTC AAGCTGGAGT GCAGTGGCAC AATCTCAGCT CACTGCAACC
158921   TCTGCCTTCT GGGTTCAAGC AATTCTGCCT AAGCCTCCAG AGTAGCTAGG ACTGCAAGTG
158981   CACACCACCA CACCTGGCTA ATTTTTGTAT TTTTAGTAGA GATAGGGTTT CACAATGCTG
159041   GCCAGGCTGG TCTCAAACTC CTGCCCTCAA GTGATCCTCC TGCCTCGGCC TCCCAATGTG
159101   CTGGGATTAC AGGCGTAAGC CACTGTACCC GGCCTCCTCC TTTAATAGAC AGGGTCTAGC
159161   TCTGTTGCCC AGGCTGGGTA CAGTGGCGTG ATCATAGCTT ACTGCAGCCT CGAACTCCTG
159221   GGCTCAGGAG ATCCTCCTGC CCTAGTCTCC CCAGTAGCTG GAACTACAGG CATAGCACAC
159281   GGGGCTAATA AAATTAATTA GGTGATAAAA TTCACTGCCC ACTGATGACT AAGCTCTTTG
159341   GACATAAAAG ACACAGACCT TGAAGGAAAA TGTGTCTACT TAATTTTGAA ACCCTATTTA
159401   TCAAAAAACA GGATGAAAAT GCAAAATGCC ATCCACATGC CAGAAGATAT CAGCTATAAT
159461   AAGTTCCCAT AAATCAATAA GGAAAAGAAC CCAATAAAAA TTATTAAACC ACAGTAAATC
159521   ATGGGTAAAT CACAGAGGCC TGAAGGGCTA ATGGACATAC AAAAAGAATC TCAATCTCAC
159581   TAGTGAAATC AGAAAAGCAC AAATTAAGTA CACAATTAGG TACCATTTTA AATCTGTAAG
159641   ACTGTCAAAA TCATAAATTA TATAAGTAAA GACTCAGGGA GTTTTGGAGG AGTGAGAGCT
159701   CTTATATTGC TTGTGGGGTA GAATTGGAAC AATTTCAAGA TCTGTAGTAT CTGGTAAAAT
159761   TATGATATGC ATCCCTCACA CCAGCATGTC ACTCCAAGGT ATCTCCCTGG AGGGAACATT
159821   TACGGGACAC AAGGAAGCAT GGATAAGAAT GTTCACAGTA GTATTGTCTG CAACAGCAAC
159881   AACAACAAAA AAACCCAACT ACACACAACT TCAATGCCCA GTCCACAAGG CAATGGATTA
159941   AATAAACTTC AGGCCGGAGA TGGTGGTTCA TGCCTGTAAT CCCAACACTT TAGAAGGCCG
160001   AGGCGAGAGG ACTGCTTGAG CCCAGGAGTT CAAGACCAGC CTGAACAAAA TAAAGAGATA
160061   GTGTTTCTAC AAAAAATTTT TAAAAATTA GCCAGACGTG GCAGTGCTTG CCTGTGGTCC
160121   CAGCTACTGG GGAAGCTGAC GTGGGAGGAT TGCTTAAGCC CAGGAATTTA AGGCTGCAGG
160181   GAGCCATGAT GGGGCCATTG CACTCCAGCC TGGGTGACAG AGTGAGACCC TGTCTAAAAG
160241   AGATAAGTAA ATAACAACTT TGCATTTTCT GCCACATTGC AAAATGGTGA GAGAGTGGTT
160301   TCTAGACTCT AGACTCTTTC TATGACTACC TTCTAGTTAT GAGATCCTAC AACACTCACC
160361   TAACCTCTCT GTGTCATATT TCCTCCTCTA TAAAGCAAAA ATGCCCCATA TAGAGAGGAC
160421   TGTGATATAA AACAAGAACC AAGAAAGTA AAGCTTTTCT AATCTGTCAC AGACTAAAGA
160481   GTGCTCAGTA TATGTGAGTC ATTATTCCTG GTGCTGGTAG GAGTGTATGT TACAACTTTG
160541   AGTCAAGTAA TATGGTACCA TATATTAAGA TTAACAACAA CCTCGGCAAT CCCAGTTTGG
160601   GGTATGTTCC CAAAAGAAAT GAAAGCACCA GGATATAAGG ATGCATGGAC TAGAAAGTTA
160661   TTGTAGCAAC ATTGTAATAA CTAAGTTCTA AAAACAGCCT GAAGCTCCAT CAGTAGGGAT
160721   ATGGTTACAT ATATTTATTA TATTCTTATG GAATATTAGA CATAAAAAGT AACGAGTAAC
160781   ATAGAAGAGA CAGTGTATAT ATGTTACGTT TGTACAAACT TAGGGAAAGA TATAGATCAC
160841   CCTACCTAGA GAAGTCAGAT TGGAGAGGGG TGGGAAAAAC CTTGAACTTT CTCCTTATAT
160901   CCTTTATATT GTTTGACTGA TTAAAATGTA TTTGTTGCAT CTGCTTGAAG GCAATGTAAA
160961   ATAAAATAAA CATACATTTA AAAATAAAAA TAAAATTTAT TCCTATCACT TTTGTAATAA
161021   AGCTGGGCAC AGTGACTAAC ACTTGTAATC CTAGCACTTT GGGAGGCAGA GACAGGCAGA
161081   TCACCTGAGG TCAGGGGTTT GAGACCAGCC TGGCCAACAT TGTGAAACCC CATCTCTACT
161141   AAAAATACAA AAATCAGCCA GGCATAGTGG TGCGTACCTG TAATCCCACG CTACCCGGGA
161201   GGCTGAGGCG CTGGAACCCA GGAGGCAGAG GCTGCAGTGA GCTGAGATTG CGGCACTGCA
161261   AGCCAGCCTG GGTAACAGCG AGACTCCATC TCAAAAAAAA ATTTGAAAAA AGAAAAATTT
161321   TAATAAACAG TGTTTAAGAG GGGAGAAATA TTTAGTTAAA AGATAAGCCC ATTTAAGAAA
161381   TAGTTTCACT TGACCCGGAA GGCGGAGCTT GCAGTGAGCC GAGATCGCAC CACTGCACTC
161441   CAGCCTGGGC GACAGAGCGA GACTCTGTCT CAAAAAAAAA AAAAAGAAA GAAAGAAAGA
161501   AAGAAATAGT TTCACTTGAA CCATATTATG ATTCCTTCTG TAAAAGATGA GAGTAGGCAA
161561   ATTGACTCAG TGAAATCCCA GCAAAACTTA CACAAAGTCT TGTTCTTCCT TCCTGTCATC
161621   TGTATAGGAT GAAATACAGA GTGCTTTTGG GTTTTGTTGT TGTTTGTTGT TGTGTATTTG
161681   AGGGGAACAC AGGTCTATAA TTCCTTTTCT GAAATCCCTG GAACAAATG GGCTTTGCCA
161741   TTCAAATTAG TTTAGAAGTT ATAAAGGCAA AAAAATGCAT ATACTCTAAA GTTCAACCCC
161801   ATCATGGCCT AAGGCAGAGC CCTGTAATCA AATTCATCAA TATATCTGCA GCAAAACATT
161861   TATTCAAATT AAGTGGGATA AATAAAGACT TTTAAATAGT CTCATCTCAG TGCCGTTCAG
161921   GGTTGGCCAC TGTGGAAGAC AGACTCAAGG GTGGCCTTCT ATGATTCCTG CCTCTTGGTG
161981   TTCACACCCT CGTAAAATTC CTTGTCTTTG AGTGTGAGCA GGGCTTATGA ATTGCTTCTG
162041   ACCAATAGGA TATGGCAAAG ATGATGGGAT ATAATTTCTA TGATTACGTT TCATTATGTA
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162101   AGACTCCATC TTGCTGGCAG ATTTTCTCTA AAGAGTCTGT CTCCTGAGCT CTCTCTGAAG
162161   AAATAACTGG CCATGTTAGA AGCCCATGTG CAAAGAGCTG AGGGGTGGCC TGTAGAAGCT
162221   GTGGGCAACC TCCAGCCAAC AGCCAGAAAT AACCAGGGCC AAAGTCCTGC AACCATCAGG
162281   AAAGAAATTC TGCCTGCTAT CTCAGTGAGC TTGGAAGTGG ATTCTTCCTT AGCCTAGCCT
162341   CCAGATAAGA ACACAGCCTG ACCAACACCT TAACTGCAGC CTTATCAGAC CCTAAGCAGC
162401   AGGCCCAACT AAGCTGTGCC CAGATTCCTG AACCACAAAA ATTGAGATAA CATATCAGTG
162361   TTGTATTAAG GTTCTAAATT ATGGTAATTT GTTTGTACTA ATAGATAACT AATATAACCA
162421   CCAAATCATT TCAGGTTAGG CCAGATTTTT GTAGCCAAAT GAATCATGAT AAAACTTTCC
162481   ATTTTCAGGG GTTTTTTTGA TTTTGTACTT ACGGATACAA ATTTGTGAAA GTATAGTCAG
162541   CACTGATTTA AAAAATCAAG GGAGCAGGAA ACTCAGTAAA TGGTTCTAAC ATTTTGGAAT
162601   CTGTAAATTG GTTGTAACAT TTGTCATCTG TGTTATCTAA GTCAAGTTCC TAAAATATGT
162661   GAATGATAGG TTATCATACT CACCTACTTT TCTTGCATTG CTCTAAGAGT TGGCTGAGCT
162721   ATTGATAATA AACACTATGA TCAGATCTAA TACCATGATG TGCTATTATG ATCATGTGTC
162781   AGTCACAGGG CTAAGCACTT TGTACATGTT GATGCATTTA ATTTTGATGA TAACTCAATG
162841   AAGTAGGAGC TGTTAATATT TTCATTTTTC AGAGGGGGAA ACCAAGTCAC TTGGAGTAAC
162901   ATGGCTAATA AGTGAAAGAA TAAGAATTTG AAAGGTTTGC ACAGATAACC AGAATGCAAT
162961   GCTCATCACA TTCACTGAGC AGTGAATCAT ACTAACTAGA GAAAGTATGA AAGCTCTACT
163021   GAAATTAACT AAACAACCTC TCTGGCTGTG AGCCTGCCAA GGGACAGGTG GTAAACTTGG
163081   TTACTGCATA AGGCCCCTTC TATCCACAGT ATTCAGGAAT TCTTTAGTGA ACATACCTTG
163141   ATGACTCCTT AACATTTTCT TCACATCGAA GTAAAGCTTG GAAACATTGC ACATAGTATG
163201   AAGTTCCAAG GAGACAGCCT CTGATGTTTC CAGCTTCACA GCCCAACTCC TAGAATAAGC
163261   AGAGGCGAGA GATTTCTTCA GAGGTGCATT CCATTCATTT CTATATACGC ACACCCCTCC
163321   CCTCCTGCAT TCAAACAGGA CTTACCTGCT CAAAGTGTCA TTCACATTCT ATAAAGAAAC
163381   AAAAAGAAAA GGTGAGCATG GGAACATCGG TATTTCATGG GGCTTGTCAT GCAGGGCTAT
163441   TCTTCTTTGC TTTACCCGAA GAAGTAAAGA GAGTTACCCT AGTCTTAGTC TTAGATATTG
163501   ATGGATACTC AAACAAAGTA ATTCCACCA GTCTTAGGTA TTGATGGATA CCCAGATGGA
163561   ATAATTCCTA CCAGCTTCTG GGAGATTCAG CATGGCAGGA TGTTTATCAA CATTTGCATC
163621   TATTCTCATC CTTGCTGAAG TCTGAGGGCC AGGAGCTTTG TCCATGCTCC CTCTGTAAGG
163681   ACTAGCTTTT GGTGATCGGA TTTCCTTCAC AGTGAGCCCA GATTAGAGAA CACTTATCAT
163741   AAAGGTCCTT AGTGGTGAAT CTGTGCACAG CCCTGAGACT GGGCCACTGC CACTAAGATG
163801   GTGGTAGCAG GTATCACACA GTGGTAAAGC AATCATGCTA TACACTCAGC CTTACAGTAT
163861   AGTCACCAAT CCTGTTAGTT AGAACCAGAA TTAATGGCTC CAGATGTTTA TCTTCCTACA
163921   GATAAAGCTG TAGATTGTAC CATAACAGCT CTGGAGCAAG GGTTCTACAA GCAAATCAGG
163981   GAAAAGGTTA TCACTCATTT TGGCTGCCCC ACTTCATCAC CCATCAGTCA CCTAGTGGAG
164041   TATTTCAGGA GAGAGTCAAC AACCAGGGTT CTCTGCACAT GGGCCAAGGA GGCAAACAGT
164101   GGTAAATGTT ATCCCGTGGT TTCATTGGC CAAGCTGTGT TCCCTCAGAA GTTTATTTTT
164161   CTAATTGACA TAAAGGTACC CTATAAATTA GTGAAGGCCA GCCTGATGGC ACTGATGTAC
164221   ATCTAAAAGA AACATTACTT TATCTTCCCA TGCTTCCTTA CCATTCTCCT TTAATAGCAC
164281   TATAACATAC CTTTTTTCCC TACTCCAAGT ACACAGCCTC ACCTGCAGCA ATTTCTGGGC
164341   TGAGCCCTGA CATTTTTCCT CCAGTTCCAG GATGTGGCTC TTGAGTTCAT TGCTCTTCAG
164401   CCCCAGACCA GCCTCATAGT CCCTCAGTCT ACTCAGAGTC TGTTGTTCTT CTTTCTCCAG
164461   CCTCCAGAGA TAAGACTTCT CTTCCTCATG TAGGAAACAC TGGAGATTCT TAAAGTCAGA
164521   CCGGATTTTT TGTCTCTGAA TCTGTACCTT CTCCTGGAGT CAAGAAAGTA TGGTCAAAAG
164581   GTGGAAGTAA ACCAAATGTC CATCTATGGA TGAATGGATA AACAAGAATG AAAGTCTGAC
164641   ACACGCTACT ACATGACAAG CCTTGAAGAC ATTCAAGCAA AATAAGCCAG AAACAAAAGG
164701   GCAAATATTG TAAGACTTTG CTTATACAAG GCATCTGGAG TAGTTAAGTT CATAGAGACA
164761   GAAAGTAAAA TAGTGGTTAC AAGGTGTTGG CAAGACCAGA AAATGGACAG TTATTGTTTA
164821   ATGGGTAGTG AGTTTCAGTT TAGAAGATGA AAGATGAAAC TGAGTTGCAG TTTGGAGATG
164881   GGAATGGTGA TGGTTGCACA ACAATGTAAC AATGTAAAAG CACTTAATTC TACTGAACTA
164941   TATACTTAAA AGTGGTTAAA TGCTTAAGTG TTATATATAT TTTCACACAA ACACACACAC
165001   ACACACAATC AGCCACTGGG ACATTATTTT CTCATGAGTC ACTGAAGCTG GAAGAATGTC
165061   CCCAGTTTCC TGCTGCAGAG TCATGTGTGG GAGGCAGGCA CTCAGATGTG GAAGAGGTTG
165121   CCTCAGATTC CTTATAGTCA CCCAATTAAT TTTCTTGTTC TTCAGCCAAG ACACAGGAGA
165181   AAGCTGGGTT AGGAGTGCTA GATAATTTAA TTGTGAAACT AGGGCCAAGT TCAAACACTT
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165241  TATCAGTTAC AAGGATAAAA AGAGGTTTTT ACTTATGATT TAAGAAGTTA GATTTCTGAG
165301  TTGGAGCGAT TTTCTTGAAG TAAAAGCTTA TAATGAACAT CACCCAGACT GGATTTTAAG
165361  ACAACCAGGC TGGTAAGAGG GTCCATAATT CTTGGCAGGG GGAGCTTGA GTGTGACAGG
165421  CATTTATTAT GGTTAACTGA GAAATACTGT TCTACTACCC TAGGGTCATC TTAAGCATTC
165481  CTATGTGTAA GACTGACAGA AATCAAGTGA AACTCTCATC TGAGGAGATG TAAAGTTGCA
165541  ATTTCCATTA GTGCTGTCTA AATTAATGCA GTGGGAGTGT GTATTCAGGG CAATTTGAAT
165601  CTATGTTCTT GGATTGCAGT CTTCAAACTT GGCCCAAATA AACTCTCTAC TTATCTTAAA
165661  AAAATAAAAA TTAAAAAATA AAAATAAATT CATACAGTGT TTTGATGACT ATGATATAGA
165721  AGAAGGGTCT TTGACTTAGG ATGAGGTGGA ATTTTTGTGT AGGAGACAGG TGCAGCTTTA
165781  ACTCTTGTAT AGACGGGTTT TCATATATGT TAGTTACAAT CAAGGTCTTC CCCATTGCCC
165841  AAGATCCTAG AAATGGGGGA AGTAAGAGTG TACTCAGGAG CTCAAGAGCA ACATCCACAA
165901  ACAAAGATCA GGGTAGAGGT TAGAGAGGAC TCCTGAAAGA GAGAAAATTG GTAATCAGCT
165961  TGTGGGATTT TACTGCAAGC TAGTGAATTA TATAAATATA AAGATTGGTG CAAAAGTAAT
166021  TGTGGTTTTT GCCTTTACTT TAATGGCAAA GACCGCAATT ACTTTTGCAC AAACCTAAAT
166081  ATTTCCATAA AAGAATGTGG CTCTGATAAT GTGGAGGTTA GTCAGCCACG GAAATAATCT
166141  GAAAGTTTGT AGTTGCAAGT GTGTAGGTTG TTGCATTACT TGTGATGTAC TTATAAATCA
166201  AGTATAGGCC GGGTGCAGTG GCTCACGCCT GTAATCCCAG CACTTTGGGA GGCTGAGGTG
166261  GGTGAATCAC GAGGTCAGGA GATCAAGACC ATCCTGGCCA ACATGGTGAA ACCCCGTCTC
166321  TACTAAAATA CAAAAAATTA GCCAGGCATG GTAGCACATG CCTGTAATCC CAGCTACTCA
166381  AGAGGCTGAG GCAGGGGAAT TGCTTGAACC CGGGAGGTGG ACATTGCAGT GAGCTGAGAT
166441  CGCACCACTA CACTCCAGCA AGACTCCATC TCAAAAAATA GTAATAATTT AAAAATAAAT
166501  AAATAAATAA AGTATATTTC TTTCATCAGC TTCATGAGCT TGAGTAGTAT GAATTTCAAT
166561  CTGGAGTGAT CCTGTTTTCT AAGTGTTCAC AAAGCTTGGT TTCTGTACCT GTAAAGTTGA
166621  GAGCCAGATG CTCCACTGTG GTAAAAGTGC CAGGGTAATG AGTTGAGGCC TGCAAACCAG
166681  GTTTATTTTG AGGTATTTAA AGTTTGAGAC CCACTCGATG CTTTTTCTAG GTAAATAGTC
166741  ATACTAATTC TGCTTCTTCT GACTGAAGTA TCAGGAATCC CAGCCAACTA CAGTTTAAAG
166801  ATGGAAAGAT TGGTGCTAAA TACTCATGGA TGTAAACCTG GAACCAGGGG CATAAGTACA
166861  AATAATGGTT TCTTCCTTGG GTTTCATTTT TTCAATCTGG TTTAGTGAGA ATAAATCCTC
166921  ATTGTGCTTT TCCTCAATCA TCCCCTATGC CTAAGCTCTA GAATGGAAAA TAGCTTGAGA
166981  TCAATGAAGT CAGATTCTTA CTTTCCATTT AGTTATTCGC ATTGCTGTGG ACAGCTTCTG
167041  CTCCGTACAT CTGTCTTCAA GTTGCTTCAG TTTTGTCACA GCTTTCTGGA GCTTTTCCTG
167101  AAGGAAAAAT TTGATAAGTG AAGCCTATTC AATTTGACTC TTCATTAGGG ACCTAGGGGG
167161  AATCCCAATC TTCTAAGATA TATTTGAATA ATAGTGAATA TTTATAGAGT CCTCATTGTT
167221  TTTTGCTAGA GAGCATGCTA AAGGCTATAT GTGCAGGAAC ATACTGATCC CCTTGGCAAC
167281  CCTGAATAGT TGGTAGGATT TTAAACTTCA TTTCTGTGCT GTAGAAAATG AGACTAAGAA
167341  AGGGGTAAAA TAACTTGCCC AAAGGGCTAT GACTGCCAGG TGGTGGAGCA ACAATTGCAA
167401  TCTCATCTGC TGACCCAGAG CCTGAGCTAT GTCCACCACT AGAGTCCTGC CAGGAAAAAG
167461  TTGGATATAG AACAAGGTAA TCATCATCTA AAAGATTTTG TAAAACAACA TGCTGAACCA
167521  AGCAAAACCA ATACCAGTGT TTGGCACACA TGAAATTTTG TGTCTTATGA GTCAGGAAAA
167581  ATCAGGATGC CAGCTGGTTA TTAGAAACAG TTCATGGAAG AGGGGAATTC TGGTATCTTT
167641  TGAACAATGG TATCATGAAT CCAATTTAAA ATGATTAGT ATTCATGTCA AGCTTTTAGC
167701  TTATTCTTCA AAACAGTTTC TCATATTTCT ATTGAAAGTG ATTTGAAGCT GACCCAAATT
167761  GCTAATTGTA GTCAATGCTG AAAGAATTGT CTCCTGTCCT CTGTAAACCC AACAAGTATA
167821  CTCATTCATT CTCGAGTGTT CTCAGGAAAA GGTTCTATGT AACTGTTTTA GCAAAAGATG
167881  ACATTGTCCT TACTATATGC CAAGTGCTAT TCTATGCATT CTATATTTTA ATGTCCTCAA
167941  AGCTTATAAC CACCTCCTGT GTATGTGTTT TAGGGAGGGA GGACACTGCT ATTATCCCCA
168001  TTTACAGATG GAGAAACCAA GGTGTGAAGA CATTAAGTAA CGTGCCCAAA ATTGCCCATC
168061  TAGTAAGTGA CAAAACTCAA TTTCAACATA AGCTGGTTCC TTTTCTTACT ACTTGGTGGA
168121  AAAGTAATTC AAATGGGAAT ATGATCATCG CAGTTATTAG CTGCTCCATG GAGTTTAAGG
168181  AAGAGCTGCC ATGAGCTGAG TGGTGGTCAT GATTGACATG TCCTTAGAAG GACTTAGAGC
168241  CTTCATACAA GACCACCTCT GCCTCATGGA GGACAGAATA AGGAGCCTGA CACTGGAGAC
168301  AACATTTTCC TCAAATTTAG GCAGGACAGA GAAGGAAAAA GGACATCAGG ACTATGCCCA
168361  TTCCTCCATG CTGCCAACAG CAAAGTCCCA CCTTCCTTAA TATGCTTTCT GGCAAGAAAT
168421  CTGGATGGTA CACAAAACCT CTCCCTCTGC TTCACCTTCC ACAACCAAGC ATTTCCAAAT
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168481  CTTTGACTCT TCTTCCTGAA TCGTGCTTAA AATCTGCCCT CTCCTCCCTT TCTTATACGG
168541  ATAGTTTGAA TTTTACTCCT TGATATTCCT TTTATCATAG ACATGCCACA GTAGCTGGGC
168601  ACAGTGGTTC ATGCCTCTAA TCCCAGCATT TTGGGAGGCT GAGATGGGAG GGAGACCAGG
168661  GGTTTGAGGC CAGTATAAGC AAGAAAGGCA GACCATGTCT CTACAAAAAA TAAAAAAATT
168721  ATCCAGGTAT GGTGGGGCAT CCCTGTAGTC CTAGCTACTT GGGAGGCTGA GGTGGGAGGA
168781  TTGCTTGAGC CCCAGAAGGT TGAGGCTGCA GTGAGCCGAG ATTGCACCAT TGTACTCCAA
168841  CCTGGGATAC AGAGCAAGAC CCTACCTCAG AAAAAAAAAA AAAAAAAAAA AAAGTAGAGG
168901  TACCAGAGTG ATATTTTCAA TGTCACTGAC CCTTCATTCC CCAAATGAAA ATCCCCCAAT
168961  AGGTGTTCAA TTTTTACGTG TCCTTCAGGA GTTACTTCTA AGATGAACCA CTCTCTACCC
169021  TAAATGTCCC TCCCCACCAC CAAAACCAGG GACCTCCAGG CAGACATTTT TGATGGTTTG
169081  TTTTCTTTAC TAGACTGTAG ATACCTAAAA GGTGATGGGT CTTTCTTCCC TGTTTTCAGG
169141  CCCTACTGCA TGGCTTTACA TATTGTGGTT TTTCAAATGA TATTCATGGT GTGAAACAAG
169201  AAAAAATGCG GGTGTTTGGT TTGAGAACAA CCTGTTCTAA AGCAAAAAGA AATTCATCAT
169261  AACACAAATG GATAGAGATA AGAGTCCAAC CATCCCATTG AAGGTCAGGA TGGACAGTCT
169321  AGATAATTGA GCAAGAAATC ATCATAAACT ATTTTTCAGA AGAATGACAT GATGAAAGCT
169381  GTATTTCCAA GTCATAATGT TAGGTTTCAA GTTAAATCAT CTCAGCTCCT GGGGAGCAGG
169441  ATAAGACTTG GTACTTACCA AAGCTCCCGG GCCCACACAC TCACCTTGTA GCCCTGGCAT
169501  ACGTCTTCAA CAAGAGCTGT GGTGTGCCCT TTGTGCTGTG GTGCCCGCTC ACAGCGCCAG
169561  CAGATGAGCT GCCCCTCATC TTCGCAGAAC AGGTGGAACT GCTCTCCGTG TTCCTCACAT
169621  GACATTTCTT GATCCGTCTC TTTGAGGGCT TCAATGAGGC TTCCCAGCTG CTTGTTGGGT
169681  CGGAGGCTAT CCATATGAAA TGGAGCCCGA CACTGGGGAC AGCAGAATGT CTCCTGCCTC
169741  AGTTGCTTTT GGCTTGGGTT TTTAAAGAAG TCTGTTATAC ACAAGTGGCA GTAGCTGTGT
169801  CCACAGTTGA TGCTTACTGG GTTCGTCATC AGGCTCAGGC AGATGGAGCA GGTGGCTTCC
169861  TCCATCATCT TCTTGGTGCT GGTGGTTGAG GCCATAGCTT TTATTGAAAA GCTCCAATAT
169921  TGGCTCTAGA GATGGAGATG AAGCAGCCAG AATTTTCCAC CGTGATGAAA ATACACCTCA
169981  CCTGCACCTC TATGTGATGA GCTGGCTGCA ACTGACTTCC ATAGGTCTTG AAGGTTTTCC
170041  TTCCAACCCC TATTATCTCA TTTTGTATTG AAGAAAGAG GACCTAAAAG GAAGAAGTTG
170101  AGGCTGAGGT TGTTTGGGCC ACGTTTGAGA ACTGCAACCC AAGTGCAGAG TTTCAAGTTG
170161  CCCTCATTAG CAAGCAGTTA CAAGTGGTTG TTTAGAGGAA AAAAAGCAGT TTTAAAGCAG
170221  TTTTAAAGTT GTTTGCCAAG AATTTACATT AAAATAGCAT AAGCTTTTGA CTGGCTATAC
170281  ATTGTTCTTT GTATTACAAA TCTCGGGAAT ATGTAGGTAA TAGATGAGGC AGCCAGTCAG
170341  GAACAAAATG CTTTTAAACA TGGGGTCTTA ACTGAAGACC TATACTCCTG CCTCACTTGT
170401  CCTGATAAAT TTTGCATACC TCACATAGCT CAGACTGCTC TAAATTATTT CATTATTTTT
170461  CTTTTCTCAG TCTTCTAACT TTTTTTTTTT TTTTAATGA GACGGAGTCT CACTCTGTCA
170521  CCCAGGCTGG AGTGCAGTGA CGCTATCTCG GCTCACTGCA CCTCCGCCTC CCGGGTTCAA
170581  GCGATTCTCC TGCCTCAGCC TCCCGAGTAG TAGCTGGGTC TACAGGTGTG CACCACTACG
170641  CCCAGCTAAT TTTTGTATTT TTAGTAGAGA TGGGGTTTCA CCATGTTGGT TGGCTAGGAT
170701  GGTCTCGATC TCTCGACCTT GTGATCCACC CGCCTCAGCC TCCCAAAGTG CCAGGATTAC
170761  AGGCATGAGC CACCGTGCCC AGCCTCTTTT TCTTTTCTTA TAAGACAAGT TCTCGCTCTC
170821  TTGCCCAGGC TGTAGTGGAG GGCAGTGGCA TGACCACAGC TCACTGCAGC CTCGACCTCC
170881  TGGGTTTAAG CAATCCTCCT GCCTCACCCT GGCAGAGTGG CTGGGACTAC AGGTATGTGC
170941  CACCATGTCC AGCTAAAGTC TTCTCTCCAG AAAGAAGAAA TGCATTGGAA TTTAGAGGAT
171001  ACACAAACAT CTAGCTGTAT AGCTAATACA GTAGCCACTA TCATGAGTAG GAATTTAAAT
171061  TTAACTTAAT AAAAATTAAA ATGAAAAAAT TCAGTTTTTC TGTTCCAGTT GCCACATTTT
171121  GATTGCTTAA TAGTTGCATG TGACTAGTGG CTACATAACA GCCTCAATAT ACAACATTCT
171181  GTTATCACAG AAAGTTACCT TGGACCAAGT GCTGGGAGAA GCAATGCAGG CTTCCTCACA
171241  AAAGCTGTAA AAGAGAGAAC TCAGGGAGTG TGAAACTCTT TCCTATTCTA GTTAACTTCA
171301  AGAATAATTG TTACCAGGCC AGCACGGTGG CTCACGCCTG TAATCCTAGC ACTTTGGGAA
171361  GCCGAGGCGG GCAGATCACC TGAGGTCAGG AGTTTGAGAC CAGCCTGACC AACATGGCAA
171421  AACCTCATCT CTACTAAAAA TACAAAAAGT TAGCTAGATG TGGTGGTGCA CACCTGTAAT
171481  CCCAGCTGCT CAGGAGGCTG AGGAAGGAGA ATGACTTGAG CTCCGGAGGG GGAGGTTGCA
171541  GTGAGCCCAG ATTACACCAC TGCACTCCAG CCTGGGTGAA AGAGCGAGAA TCTGTCTTAA
171601  AAAAAAAAAA AAAAGAATAA TTGGTACCAG AATTACTCTT TGTAATTAGT AGTAACACTT
171661  ATGCAATTGG GTGATCTGTG ACAGATTCCA TTGAAGGAGT ATGGGGAGCT TCACCCCAAT
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171721  ATATGACTCC CTGGTATAAT GAGTATTTTG AATTAAAGGC CCTTAGAGAT CAGCAGATGC
171781  TGGAAGAGAC TTTTCCCCTA TCTACATAAA GACCAGTCAC ACTAGACAAG AAGAACAATT
171841  GTTTTTCCTT CCAACCCCTA TTATCTCATT TTGTACTGAA GAAAAGAGGA CTAAGAATGT
171901  AACCAGACCT AATCAGACAC TTTCACAAAA TAATGTCTGT CTCTCAGGCT CATTCATTTT
171961  CCAAAGAGAA CCATTTACAA GTTAAACTCT GTTCCTCCAT TCATTCATCC TCCCAAATAT
172021  TCATTTATTC TCCCTAGTAA TCATTTACTG CCCCTCAAAG AATTACCTAT ATTCTCCTGA
172081  TATCACCCTT CCCCTCTGAA ATAAATATGT ATACATGTAT AAACGTTATA CATACATATT
172141  TATACAGTAT ACATACATAT TTATACATAC ATACATATGC ATACATATTT ATATTTATGT
172201  ATTTATACAT AAGTATTTAT AAATAAGGCT ATATAAGTAT CTACCCCCAT TGGCAGAGGG
172261  GGTAATCACT CTGTGATTCT AGCCCATGTA CTTGTTAATA AATTTGTATG CCTTTTCTCC
172321  AATTAGCCTG CCTTTTGTGA GTCGATTTTT CAGTGAACTT CAGAAGGCAA AGGGGAAGTG
172381  TTCCCTTGGC TCCTACACCA TCATGACAAT AAAATTTGAC TCCACCTCGA CCCCCCCCAT
172441  CCCCCACAAA GAACAACAAC CAACACTGGT AATAAGGTC GGTTGTTTTT TGTTTGTGTT
172501  TTTGTTGTTG TTGTTGTTGT TGTTGTTTTT GCTTTCAGGA GCAGAGGTAT AATAGGCAAA
172561  AGAAAGAGAA AGGAGAATAG TGAATACCTC TTCTGCAGAG AGGGGTGCCT AAGTGGGACT
172621  TCCCTGGCTA ATAACGTCTT GCTAGAGACC CAACCAGGAG GATAATGGAA GCAATCAAGG
172681  CAACCAGAAC AACCAGAAGA ACCAGTTTAT CCTTTTTGTG CCCTCTCCCT AAACTGAGGG
172741  AATAAGAATT GGAAAGAAGG CTGCAGAGCA GAGGGTTTGC TCCTGAGGAG CAGTTATTTC
172801  TATGGGATCA GAGCTCCTGC AGAACTGGGG AGTTTACTTT TACTATCTCT TCTCCAGGAC
172861  AGGACCTATC TCAAGAGACA TGTTCAGAGT GATTGCAACA TAAAGAGTTT GCAGACCCAA
172921  GGAGGTAGGG AAGGCAGAAA GAAGATGGGG GAGGCCAGGG ATAGGCAACA GAGGAGTGAC
172981  CAGGAGCGAA AAAGCCTGCC TCTTCTGAGA ACCTAGCTGG GCTCTCCCTG TACCCCCGAT
173041  CCCTCCCCCC CGCCCGCCCC CACACCCCTA CTCCTGGGAG CTCCTCTAGG ACAGGGGCAG
173101  AGTCAGGAGG AAGTTTGAAG AGTGCCTAGA ATAAAAAACA GTAATTTAAC TACAATTACC
173161  GGGTAGGCTG TTTTCCTCTC ACAATTTGAT CAGTCTCTTG AAGCCACACA GAATTTCTTC
173221  TGAAGACGTG TATTCCTTGG CAGGCTATTT CCTCCAGTGA TACACCAGGC CCCTCTCTGC
173281  TGGGGTCACT GCTCTTCTGG GGAGATGGGG CTCCCCTCCT TCCAAGGCTC CAGGGTTCCT
173341  GTCCTGGGCC CCACTCATCT AAGTTCTGAA TCTTCTGAGA TTTGGTGTAA AGTCTGGTGA
173401  AAGAAAGAGC AGGAAAGAGG TGAGAGCTGT AAAACAAAGA AAGTCCTGAC CATTTTCAGA
173461  GTTGGAGGGG CCCTGCTGTC ACGAAATATA TTCCCCACCC CACTTGCCAT CAGTACACAC
173521  TCACATATCC ACTGAGAAAA CCTTAGCCTG GACCTTTTCC GTAACCTTCA CTGCTCAGAC
173581  ACTTACATAT TCGCTGCTAG TCCCCTCTGT TGCTGCCACT TCCTGGGTCA GGAAGTTAAC
173641  TCAGACCGGA TTAAACTGAG AAGTGAAACT ACTGTGGGAG GCGGGGCTCA TAAGATTTAG
173701  GAGAAACTA GTGACGTTGT TCATATCATT TGCACTCCGC CTCTCCGGTA AAGGAGGGGG
173761  AAACGTAGGA AGAAAATATC CTTCTTTTAC AGCAATAAAA AGAAGGAACC AATTAATAAC
173821  CCTGTAAACT ATCATGTGAC CCCAACACAG AGTATCTAAA AACAGGAAGC CTGCAGAGGT
173881  TCAGTTCACA GACTCTGATT TGAGATCTTT CTACTTTTGC CACCAACTCC CTTGGGAGTC
173941  CTTAAGCCTT CCTAGCTGAT GTTACTTCTT TTGCTATTTA TGGGTTGCTT GTGGTTCTAT
174001  AACTGCTCTG AAGGGTGTGG TGGAAAAAGG GGTGGTAACA GCAGTAGGAC TCATTGGCAT
174061  CACAAAATTC ATCTGAGTCA GCTTTCTATT CTTCTCTGTC CCGTTCTGTG TCTTGTTTTT
174121  CTCCTTGCTG TCCTTCTGCA GGACTCAGAT CTTCTTCAAT AGCGAGGGTC AGCCAGGATA
174181  GAAAATGGGA GTCACTAGTG GCCCAGCAGT GAGTGCCCCC AGCTTAGAGC TGTGTGGGAT
174241  CCCTGGGACC ATCACTCTGC TTTGTGCTTT GTGGAGAAAA GGCTGTGGGG TCCAGGGTCA
174301  AGTCCTTAAT GACTTAGCTC CAGCTTCTCC ACTTCAAAAT GAAAGGAAAA GTACTATCAC
174361  CACCCGTTAG AATTATTATT TCATGGGGAA AAAAGATGGA TTACTATCTC ACAATAAGAG
174421  CTTGTCACAT TTATAAGTCT CAGGTGTAAG AGGCATTTAT GATAACAACA TAATAAATGC
174481  TGGCTTAAGT AGATGCAGTG GTCCAAGGGA ACCAGTAAGG GGAGCTCAGG ACACAGGTGG
174541  GAGGAGAAAT TAAACTTGAA TTCTGGGAGC CACTGGCCTG TCTGGGCCCC TGGCCTGCCT
174601  GCTGACCCTG ATAGCCAATG GAACATGGAG TTTGGCCCAG CTGCAATCCC TCTGGTCCAA
174661  CTACTCAAAA TAAAGGCAAG ATTGGGAAAC ACGTTCCTTT CTTCCTATAC CAAGCAGAAG
174721  ACTCTTCAGC ACTGCACCCT CCTGGGTGCT CACAGAGCCT TCTGTTGTTT TGCCACCTAC
174781  GATTCATCAT GCCCTGGCAT GATGGTTGCA GACCCCATGC ATAGCATGGG ACATTCTACT
174841  CCTGAGGCAA CCAGCACACA GAGAGAGGAG AAAGAATGAG CCCCTGAATC CTTGGTCCCA
174901  CGATGAGTCC TTGCAGATAT CTACAACTTT CATTGTTGTG GATGTGACTC TGTACCCAGG
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174961  CATGGCTCAT  TCCAGATCTG  TCCTATTGTC  AGAGGTGTTC  AAACCAGAAT  GACTCCATTT
175021  TGAATGGGGG  CTAGGTAAAA  TAAGGCTGAG  ACCTACTGGG  CTGCATTCCC  AGGAAGTTAG
175081  GCATTGTAAG  TCACAGGATG  AAATAGGCAG  TTGGCACAAG  ACACAGGTCA  TAAAGATCTT
175141  GCTGATAAAA  CAGGTTGCAG  TAAAGAAGCT  GACCAAAACC  CACCAAAATC  AAGATGGCAA
175201  CAAGAGTGGC  CTCTAGTCAT  TCTCATTGCT  CATTATACAC  GAATTATAAT  GTGTTAGCAA
175261  GTTAGAAGGC  ATTCCCACCA  GCTCCATAGT  GGTTATAAA  TACCATGGCG  ATGTCAGGAA
175321  GCTACCCTAT  ATAGTCTAAA  AAGGGGAGGA  ACGCTTGGTT  CTGGGAATTG  CCCACATCTT
175381  TCCCAGAAAA  CATATGAATA  ATCCACTCCT  TGTTAGTAC  ATAATCAAGA  AATAACTGTA
175441  AGTATCTGTA  TTAGTCCATT  TTCACACTGC  TGATCCAGAC  ATACCTGAGA  CTGAGTAATT
175501  TATACCAGGA  AAAAATGTTT  CATGCTCTTA  CAGTCCCACG  TGTCTGGGA  GACCTCACAA
175561  CCACAGCAGA  AGGCAAGGAG  GAGCAAGTCA  GGTCTTACAT  GGATGGCAGC  AGGCAAAGAG
175621  CTTGTGCAGG  GAAATTCCTT  CCTATAAAAC  CATCAGGTCT  CATGAAACTT  ATTGACTATC
175681  ATGAGAACAG  CAGTATAAAT  TACTCAGGGA  AAGACCTGCC  CCCATGATTC  AATTACCTCC
175741  CACCAGGTCC  CTCCCACAAT  ATGTGGGAAT  TTAAGATGAG  AGTTAGGTGG  GGACACAGCC
175801  AAACCATATC  AGTATCCTTA  GTCCAGAAGC  TGATGCTCTG  CCTGTAGAGT  AGCCATTCTT
175861  TTATTCCTTT  ACTTTCTTGC  TTTCACTTTA  CTGTGTAGAC  TTGCCCCAAA  TTCTTTCTCA
175921  CACGAGATCT  AAGAACCTTC  TCTTAGGGTC  TGGGTTGGGA  CCCCCTTTCT  GGTAACACTA
175981  TCAAAGGATC  AGGAAAAGGA  AGCTAGTGAA  TGCTAAAAAG  GAAACAAACT  ACCATTACCA
176041  ATAATAACAG  CAAGACAAAA  GCAAAACGGA  TTGTGACAGC  TGTCCCATCT  CACACCTGTT
176101  TCCCATTGCA  GGAAGGAGGG  GCTGGTTCAT  GCACAGAGTG  GCCAATATTA  GAAGCAGAGA
176161  GGGGGTGCAG  ATGAGACTTC  AGGAATATGT  TGACAAAGGC  AGGCCTAGGG  AGAAATCAAC
176221  CTGAACTATC  CCCAAGGAGG  AATGCATTAT  CTCTAATATG  TAAAGTTAGG  CTTGATCCTG
176281  TGATTATGGG  ATATAGGAGT  CCAAAGACTC  ACAATGGGAA  GTAGGTCACT  AGAGTCTCCT
176341  TCAGAAGCTC  TGTACTGTGT  GTTCCCACTG  TGGGCAAGAG  TCAGCACTCA  GCTATTCCTA
176401  GAATGCCTTT  CCTCAACTCC  TTCAGATTTT  GCCTCTCAAC  TAACCCTATC  CTGACCACTT
176461  GTTAGCAAGT  GTACCCCTCT  CTCCCTCCCA  AACATTTTCA  AATCTATTTT  GTTCCCATGG
176521  CACTTATCAC  TGAATATTTT  ACTAATTTAT  TTTGTTAGT  GTTTGCTTCC  CTCATGAGAA
176581  TGCAAAGGGA  TGGATTTTTT  TCAATATTGT  TCACTGATGA  ATCCCAGTAA  CTAGAATATT
176641  TCTAAGCATA  GTGATGTGCA  TTAAATCAAA  GAGTAACTTT  CTGAATTGCA  CTAAACACAC
176701  ATCACAAGAG  GTGTGTGCAC  ATATGTGCAT  GATGCACGTA  GTGTGGTGTG  GGTGTTGTGT
176761  GGGGTATGTG  GTACTGTGTG  TGCTGTGTGT  GGTATGTGAT  ACATAGTTTG  TGTTAGTGTG
176821  ATGCATGTGA  TGTGGTATGT  GTGTGCGTGT  CCATACATAT  TAGGGGTGGC  GGGGATGTTA
176881  ATATGTCAAA  TGGTACTAGA  AAGTATCAGA  ACTCATGGTG  CTTACTGGTT  TCCCAGAGAG
176941  CTGCTTCTCT  CCCACCTGTA  GGATATACTG  ATGGTTTGGA  CAGAGAAGAA  ATAAAAAGAA
177001  GGCTGTGACC  TACTGGGCTG  AGGAAATAAA  AACGAAAGTA  AAAGAAGAGC  TGGGAAAAGA
177061  GAGTGGAGGG  GCCAAGGGAA  ATTTCCCCTT  TGGCTTCTGG  GGAAACTTTG  CTGAAAAATC
177121  AACTCACAAA  TTTATTAACA  TGTACACAGG  GAGAACCATA  GAATGATTAT  CCACTTCCCA
177181  AGAGGGCTTA  AAAGCTTATA  TATTATCCTG  GCAAACAGA  TTATGGGAGG  GGAAGAAGAG
177241  AAACTCTGTT  GATGGGATTA  CTGTTGCGGA  TTTTTGCTCC  TTCGCTCAGC  TAGGTCCGGG
177301  TTTTTGTCTC  ACAGCCAGGA  AGAATTAGGC  ATGCAGCCAT  CAAAGAATGA  GTGGAGTAGA
177361  ATTTATTAAG  TGAAAGGAAA  GCTCTCAGCA  AAGACAAGGG  TCCTGAAAGC  AGATTTCTGG
177421  TTTGCTCTTC  ACAGTTGAAT  ACTAGGGCTT  AAGACTCAAA  TTCCTGACAA  CTCCACCCTG
177481  TCCTACCAGT  GCATGCAGGC  CTTTAGACTG  AGCTACTCCA  TATTGATTAA  TTTCCTGAAC
177541  TGCGCATGTG  TTAAGGAAAG  GAATCATCCA  CTGCAGGCAT  GTTAGGCAA  GCCCCCTGTG
177601  CAAGTTCCCT  TATCTGCACA  AAACATCCGG  TGTAAGCACT  TGTGGGGCAG  GTCAGAGGTT
177661  CTCTGGGTAC  CATTCCCTTA  CTGTCTGCCT  AAAGCAAGCT  GGCCAACTCC  TTTCATTACT
177721  AGGGAGAGTA  AGTAGATCAG  GGAACAGAGA  TTAACTTGAA  CATTATCTTG  TGAAAGTCCG
177781  TTCGGGCATG  GTTACATTCT  TGGTCTTACA  GGAAGGGTAA  ATAAAAATAA  TTGCTCTTTT
177841  TGGTGGGTCT  GGATCTTAGG  TAGATAAAGA  AACTTTAATT  CCACGATGTG  TTTTGGTAGG
177901  GATAGTTGGT  GGCAGGGATG  TCAGAGAGAC  TTTGAGGCTT  CTTCAGTTCA  ATATGACCAA
177961  GGGCCATATA  TTAGGGTATC  AATTTCTGAG  CCCCAACAAG  AGCTTAGGAG  AGATGTGATA
178021  GCATCACAGT  GTGAAAGCAA  TTTTTTGTCT  GTTTTTAGAG  ACAGGCTCTT  GCACTGTCAC
178081  CCTGGCTGAA  GTACAATGGT  ACGATCACAG  CTCACTGTAA  TCTTGAACTG  GGTTCAAATG
178141  ATCCTCCCAT  CTAAGCATTT  CAAAGTGTTG  GGATTACAGG  CATGAGCCAC  GGTACCCAGC
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178201   CTGAAACTGC ACCCACTTTC TGATAAACTT TTCAAATGAC TAAAGGGGAG AGAGTAAGCA
178261   CTACTCAGAG GTAGGAAGAA AGGACACAGG ATTATAGGAT TAAAACAACA ACCACCAAAA
178321   AAAACCAGAC CGGTGTGGTG GCTCACACCT GTAATCACAG CACTTGGGGA GGCTGAGGTG
178381   GGGGGAGTCA CTGGAGGCCA GGAGTTCGAG ACCAGCCTGG CCAACATAGC AAGACGCTGT
178441   CTCTATTAAA AAAAAAAAAT ACCTGCCTTG AGCTAATCAG AATCATGGAC CCTGACAAAG
178501   GATGTCCCAA AGTAAGTCTT AGCATTTTTT TTTTTTTTTT GAGACAGTCT CGCTGTGTTG
178561   CCCAGGCTGA AGTTCAGTGG CGTGATCTCG GCTCACTGCA ACAGCTGCCT CCCAGGCTCA
178621   AGCAATTCTC CCTGCCTTCA GCCTCCCAAG TAGCTGGGAT TACAGATGCC CACCACCACG
178681   CCTGGCTAAT TTTTGTTTTT TTTAATAGAG ATGGGGTTTT GCCATGTTAA CCAGGCTGGT
178741   CTTGAACTCC TGACCTCAAG TGATCTGCCC ACCTTGGCCC CTCCATAGTG CTGGGATTAC
178801   AGGCGTGAGT CACTGCACCC GGCAAAGTCT TAGCATTCTT TACAAACAGT TTGTACCCGT
178861   ATCTCTAAAA GGGAGTAGTG AATTTCACCC CAAAATATGG CTTCCTGATA TAATGAGTAT
178921   TTTGAATGAA AAACTCTTAG AGATCAACAG ACACTAAAGA GACTTTTCCC TAGGTACATA
178981   AAAATAGGAT GGCCCCACCA GCGAGAACAA TTGTTCTTTT CTCCCTCCCT GTTATCTCAT
179041   TGTGCATTAT AGGAAAGACC AAGAATGTAA CCACACCTGA ACAGACCCTT TTATAAGATA
179101   ATCAGTCTCT AAGCATCATT TAAATTCCAA GGAGAACTAT TTACAAATTT ATCTGTTCTT
179161   TGATCCAATT AGTCTCTCCT GGTAGTTACA TATTGCCCCT CAACAGAATT CCTCTTCTTC
179221   TGTTTCCCAT AACCTATTTT GCAAGGATCA AGCCCTGTT ACTTCTTCAA CTTCAAGTTG
179281   GCATATAAGC TTCTAAATTC CACTGGGATA TTGGTACTAT GTGCATGAGG AGAACCACAG
179341   AGTAATTAAA TTGTAAAGCC TTTTATCTTA TGAATCTGCC TTTTTTTGTG TTCATTTTTC
179401   AGCAAAACTT CCAAGGGCAA AGGTATAAAA CAAAAATAAA ATTCTAAAGC CCCCCAACCA
179461   TCTGAATAGA CTTTCTCTTC AGTCAGGCTT CTTAAAATGT AACCTGAAAG ACTGGCTCAG
179521   GCCATTAAGG GAAGTGGGGG TTGAACATGC CTCATTATTC CTCTCTGGCA TTAACATCAA
179581   CACAGCTTTT AAGTCTGATA AGAAACATTT TACAACCTAT TCTCTCTGAA GCCTGCTAGC
179641   TAAAAACTTC ATCCCATAGT ACAACTTTGG TCTTCACAAC CTGTTATCAC AACCTAGTGC
179701   TCCTTTCTAT TAATCCCAAA TCTTTATACA AACTCAACCA ATTGTCATCA CCTCCACCCC
179761   ACTCCTCCGC TGCTTCCAGT TGTCCCGCCT CTCTGGACCA AACCAGTGTA CATTTCTTAA
179821   ACGTATTTGA TTGATGTCCC ATGCCTCCCT AAAATGTATA AAGCCAAGGT GCATCCCAAC
179881   CACCTTGAGC GCTTGTTCTC AGGACCTCCT GAGGGCTGTG TCATGGGCCA TGGTCACTCA
179941   AATTTGGCTC AGAATAAATC TCTTCAAATG TTTTACAGAG TTTGGCTCTT GTCATGACAC
180001   AGATGACTGC TTCACTGAAG CCTGCTCTGG AAGTGAGTGG GGGTTTTGCA AGGATAATTT
180061   TCCCCGGATA GCCCCAGAAG CAGCTAGTAA TAATACACTT AAAGGTAGCT AAAATGCATT
180121   GAACACTTGT TTTGTGCCAG ACCTATGTCA ACATTTGCTT TGTGCCAGGC TTATGCCAGT
180181   ACTCCTGATT TGTTAATACA TTCTAAATAA AAATTCTGGA GTTTCAAATA TAATAACTGA
180241   AAAACAGAAA ATAAATAAAA ATATATAATA ACTGAAATAA AAATTACTA AGGCTGGGGA
180301   TGGTGGCTCA CTCACACCTG TAATCCTGTT ACCGGAAAGG GGTCCGTCCA GATCCAGACC
180361   CCAAGAGAGG GTTCTTGGAT CTCACACAAG AAAGAATTCG GGCGAGTCTG TAAAGTGAAA
180421   GCAAGTTTAT TAAGAAAGTA GAGGAATAAA AGAACGGCTA CTCCATAGGC AGAGCAGCTC
180481   TGAGGGCTGC TGGTCGCCCA TTTTTATGGT TATTTCTTGA TTATGTGCTA AACAAGGGGT
180541   GGATAATTCA TGCCTCCATT TTTTAGACCA TATAAAGTAA CTTCCTGACG TTGCCATGGC
180601   ATTCGTAAAC TGTCGTGGCG CTGGTATGAG CATAGCAGTG AGGACGACCA GAGGTCACTC
180661   TCATCGCCAT CTTGGATTTG GTGGGGAGCA GTGAGGATGA CCAGAGGTCA CTCTCATCGC
180721   CATCTTGGAT TTGGTGGGGT TTAGCCAGCT TCTTTACTTT TTTCCTTTTT TTTTTTTTTT
180781   TTTTTTTTT GCCCAGGCTG GAGTGCAGTG GCACGATCTC AGCTCACTGA AACCTCCAAT
180841   TTCTGAGTTC AAGCGATTCT CGTGCCTCAG CCTCCCAAGT AGCTGGGATT ACAGGCATGT
180901   GCCACCACAC CCAGCTAATT TTTTATATTT TTAATAGAGA CCGGGTTTCG CCATGTTGCC
180961   TACGCTGATC TCCAACTCCT GCGCTCAAGC CATCCAGCCA CCTTAGCCTC CCAAAGTGCT
181021   GGGCTTATAG GTGTGAGCCA CCCCACCTGG CCTAGCCGGC TTCTTTACTG CAACCTGTTT
181081   TATCAGCAAG GTCTTTATGA CCTGTATTTT GTGCCCACTG CCTGCCTCAT CCTGTGGCTT
181141   ACAATGCCTA ACTTACAGGG AATGCAGCCC AGCAGGACTC AGCCTTATTT CACCCAGCTC
181201   CTATTCAAGA TGGAGTCTTT CTTGTTCAAA TACCTCTGAC AAGCCCAACA CTTTGGGAGG
181261   ATGACACAGG AGGATTGCTT TAGCCTAGGA GCTCAAGACC AGCCTGGGCA ACACAGTGAG
181321   ACCCCATCTC TAAAAAAAAA AAATACAAAA AAATTAGCCA GGCATGATGG TGTGTGCCTG
181381   TAGTCCCTGC TACTCAGGAG GCTGAAGTGG GAAGATGGCT TCAGCCCAGG AATTCAAGGC
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181441   TGCATTGTCA GAGGCATTTG AACCAGAATG ACTCTATCTT GAATAGGGGC TGGATAAAAT
181501   AAGGCTGAGA CCTGCTAGGC TGCATTTCCA GTATGTTAGG CATTCTTAGT CACAGGATGA
181561   GATAGGAAGT CAGCACAAGG TACACATCAC AAAGACCTTG CTGATAAAAT AGGTTGTGGT
181621   AAAGAAGTTG GCCAAAACCC ATCAAAACCA ACATGGCCAC CAAAGGGACC TCTGGTTGTC
181681   TTCACTGCTC ATTATATGTT AATTATAATG TATTAACATG CTAAAAGACA CTCCTACCAG
181741   CATCATGACA GCTTACAAAT ACTGCGGCAA TATCTGGACT TTACCTTATA TGGTCTAAAA
181801   GGTGGAGGAA CCCTCAATTT TGGGAATTGT CCACCCCTTT TTTGGAATGC TCATGAATAA
181861   TCCACCCCTT GTTAGCACA TAATCCAGAA ATAACTATAA GTATGCTTAT TTGAGCAGAC
181921   CACGCTGCTG TTCTGCCTAC AGAGTAGCCA TTCTTTTATT TCCTTACTTT CTTAATAAAC
181981   CTGCTTTCAC TTTACTGTAT GGACTTGCCC TAAATTCTTT CTTGTGTGAG ATCCAAGAAC
182041   CCTCTCTTGG GGTCTGGATC AAGACCCCTT TCTGGTAACA TCTTTCTGGT GACCACGAAG
182101   GGACAATACT GAGGAGACTC TGAAGCCAAA GGAAACAGAC TACAGCACCA ACTGGCTGAC
182161   TTTGGGTAAG TGGTGGAGTC CCCGGGTAAA GGATAGGATT GGGTTAGAGG TGCAACTTAG
182221   GGGAGATAGG GTCTCTCCTA AGACAGAGAG CGTTTCAGTC CGCTCTTAAT AAAGGGCAAG
182281   AATGCTTGAC CGAACTTGGG TTTGAGACCC AACTTAGGAA GGCTACAGTC CTTAAGATTT
182341   AAGGGGTTAG AGGCCCCTCT CAGTAAAGTC TCTCTTGGTT AAAAACGGAT TTAGCATTAG
182401   GGGATGTTAA CTGCTATTCT GTTTGTATTA ATCTTCCCTG TGCTCTTTGC TGACAGCTAT
182461   GGGTGACAGG ATTAGGCATG TACAGGATCA CGGGACATTG GGAACTTTTC TTCTCTCCAA
182521   AAGGGGAAGC TTGACAGCTG ATAGGACTGT TGGAAAGAT CCCTTTGCTA TGACAAGCAG
182581   CCGCCTGAAC TTTTGATTCA GTGTTGCTGC AATGGGTGGG TCTTTCTCTG GCCTCTGTGA
182641   ACTCCTCACC TTCCCCACCT CACCACAGGC AATGCTTTTC TCCCTTTCTC TCTTTTCTCT
182701   TTTCTGTCTT TTCTGTTACT TGAGACAACC ATCTTGCCCA GAGACCATAT GTTGAAACTC
182761   CTGGTCAGAA GTTTGATTAA AGATGAAAGG GCCTATCTGG GGGCAAGTTT GAGCCTTCCC
182821   AGTTAGATAT TGGGTGCTAA GTGGAGTGGC CAATGTCTAT GTTTTGTCAC ATGTATATTG
182881   CTCTGGCTGA AATGGAAAAC GTTAATTTGG TTACTTTATG TGGCCATTGG GCAGCATCTT
182941   ACAAAAGTGA GAGACATTTA TTTGCCTGTG GTTCCATGAA ACAGAAAAAA GTTGGTTTTC
183001   CTTTGTGTCG TAGCTTGGAC CCAAGGGCTT TGCAGTGAGC AAGGTTGCTA GCGCTGCTCA
183061   GTGAAAGAGA ACCCAGAAAC CTGGCATGCC AGCAAAGGG TAAAGATTTC TTACCAGTCA
183121   GGCTTCTGGC CTCTCTCTCT TAGTGAAAAC TGAATGAATG GTAAAAATCA CTGTTTATCA
183181   CCTCTGTAAA GTTTTGATTA ATGGGAACAA GGATTGTGG GGCTAGTCTT AAGCTGTAAT
183241   GAATCTGGTA TACTTTGTGA TATCAATTTG TCTTTCTGTA TTACTCTGTC ATAAAGAGGA
183301   ATATGGTAGG ATAGAACATG GGCTTAGGAC TCCATAAGCC TGCTGTTCAA GCCAGCCCAG
183361   TAAACTGGTC CGTTGCAAAG TTTATTACAG GTCCCTGGAA AAAAAAAAAA TTAAAAACTG
183421   GATGAAGTTT CCTTCTCATC TTGTTTTATG TCCTTTGGAG CTTCACCTTG TAACCACGTG
183481   GCGGTACTTT CTCTTGGTCT CTGCCATCCA GGGAACAGGA ATTTTGGGGT TTATGTAATA
183541   GTTAACTCTA AAAATTATCT CAAGCCATTG CAAGCTCAAA ATTGGCTGCT CTGGACCCCT
183601   TCTGGGAAGG GCAATGGAAA CTAACCAGTG TTGTAGCTCA GCAGCTAAGG ATTTGTCATT
183661   TTATAATGGC GGCCAAGGTT CAATCCTGGC TTAGGGAATG AGTACTTTCT GATTGATATC
183721   TGTGTGACCT TTACCATTTG TTGATTCTGT TCTCTTCCCC TCCACACACT GTCTTGAGTT
183781   TTCCTCTCTC TGAGAACCTG GGAGATTATC TTTGGTAAAG TTCAAAAGCC AGAAATAATG
183841   GCCGTGTGGG ATGGCTAAAG TTGAGTAATA AGAAACTTAA AAGGACTCCT TTTTTTTTTG
183901   CTTTAGAGTG CTATGGTTTA TGGTTAAAAG CTTAATTAAA AGTGGATATT CAATCTCTAA
183961   AAGCCTGGGA CTCCTTGGGA AAAGCAGAGG AGGCACCACA GACCCCATTT TGGGAAAACC
184021   TCTGTTTTCC TCATGAAACC CCAGGAACTG GAAGTGGATA GATCCTTCGC AAAATCTAAG
184081   GCTCTGTTTG GCTTTGCATT ATGTTATCTG ATGTTTTTGA CTTTTGGGGG TATCAGAAAT
184141   TACTTTGCAT TATGAGGGAG ATCTGGTGTG TAATAACCAG GTAGGAAATA TACTTCTGGG
184201   GATAGCTAAA GGCAAATATA GGTGAATACT TGGCTATTTG CACTTTTGGA TCACAAGAAG
184261   CATTCTCTTG ACTACCTAGA AGGTATGGAA ATGTCTCCAT CCCCACCGAG AGATAAGATT
184321   CCCAGGGGAG ATGGCTGATC CCCCAAAAGA GGGCTGATTC CCTCTTTTGG GATCCAGGAT
184381   CTGGTATAAA AATGGGACCC TGGCCAGGCA CAGTGGCTCA CGCCTGTAAT CTCAACACTT
184441   TGGGAAGCCT CAGAGTTATG AATGTCTCAC CATACTGACA CTTTGTGACT GAGCTCCTCT
184501   CTACCCTGGA CACAAGAGAC CCTAATAATT AGACAGGAAT ATCATTGCCC CTATTTAGTC
184561   TGAAGAAGTT ATAGAAGATG GATCTTATC CCACTGCAAT CCTTAGGATT AAGGGTTCCC
184621   TGGTAAAAGG GAGTGGGAAA ATATGTCAGA GGCATTTGAA TCAGAGTGAC TCCATCTTGA
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184681   ATAGGGGCTG  GGTAAAATAA  GGCTGAGGCC  TGCTGGGTTA  GGTTAGGCAT  TCTAACCAGG
184741   AGTTTAGTCA  CAGGATGAGA  TAGAAGGTTG  CACAAGGTAC  CCGTCACAAA  GACCTTGCTG
184801   ATAAAATAGG  TAACGGTAAA  GAAGCCAGCT  AAAGCCCACC  AAAACCAACA  TGGCCACAAA
184861   AGTGACCTCT  TGTCATCCTC  ACTGCTCATA  TACACTAATT  ATACTGCATT  AGCATGCTAC
184921   AAGACACTCC  CACCAGTGCC  ACGACAGTTT  ACAAATACCA  TGACAACATC  TGGACGTTAC
184981   CTTATATGGT  CTAAAACGGG  GAAGAACCCT  TAGTTCTGGG  AATTGTCCAC  CTCTTTCCTG
185041   AAAAATTCTT  GAATAATCCA  TTAGTTTAGC  ACATAATCCA  GAAATAACTA  TACGTCTGCT
185101   TATTTGAGCA  GTCCATACTG  CTGCTCTGCC  TATGGAGTAG  CCATTCTTTT  CTTTTATTTT
185161   TATTTTTTAG  ATAAAGACTC  GCTCTGTCAC  TCAGGCTGGA  GTCTGGAGTG  CAGTGACGTG
185221   TTTTGGCTCA  CTGCAACCTT  CACCTCCCGG  GTTCAAGCAA  TTCTCCTGCC  TCAGCCTCCC
185281   AACTAGCTGG  GACCACAGGT  GGGTGCCACC  ATGCCTGGCT  AATTTTTGTA  TTATTAGTAG
185341   AGATGGGGTT  TCGCCATGTT  GGCCAGGCTG  GTCTCGAACT  CCTGGCCTCA  AGCGATCCAC
185401   TTGCCTTGGC  CTCCCAAAGT  GCTAAGATTA  CAGGCATTAC  CCACTATGCA  TGACCCATTC
185461   TTTTATTTCT  TAACTTTTTT  TTGTTTTTTT  GAGACAGAGT  CTCACTCTGT  CACCCAGGCT
185521   AGAGGCTGGA  GTGCAGTGGT  GCGATCTTGG  TTCACTGCAA  CCTCTGCCTC  CTGGGTTCAA
185581   GCGATTCTTC  TGCCTCAGTC  TCCTGAGGAG  CTGGGACTAC  AGACATGTGC  CACTACACCC
185641   AGCTAATTTT  GTATTTTTAG  TAGAGACAGT  GTCTTGCCAT  GTTTGTCAGG  CTTGTCTCGA
185701   ACTCCTAACC  TCAAGTGGTC  TGCCTGCCTC  AGCCTCCCAA  AGTGCTGTGA  TTACAGGCAT
185761   AAATCACTGC  GCTCGGCCCT  TCTTTACTTT  CTTAATAAAC  TTGTTTTCAC  TTTACTGTAT
185821   GGACTAGCCC  CAAATTCCTT  CTTGTGTGAG  TTCCAATAAC  CCTTTTGTGT  GTGAAAGAAT
185881   TTATGGCTGC  TGTTCAGGCT  GGAGCAAGCT  GGAGCTCATG  CTGCTGCTCA  GACTGGAGCA
185941   TGCGTGATCT  GTGATCCCAG  TAAGAGGATC  ATGGTCACTC  CAGCCTGAAC  GACAGCATGA
186001   TATCTCATCT  GTAAGAAAAA  AAAAATTACT  AGAGGGCTTT  AACAGCAAAT  TTGAGCAGCA
186061   AAAAGAAGTA  ATCAGTGAAC  TCAAAGATAG  GTCAATTGAA  ATGATCTACT  CTGAAAAACA
186121   GAAAGAAGAC  AGAATGAAGA  AAAAGAAATA  GAGCCTTAGA  GACAGGGGAT  ACCATCAAGC
186181   ATACTAATAT  ATGCATAATG  GGACTCCTAG  AAGGAGAAAA  GTGAGAGGAC  AGGGAGAGAG
186241   AATGTTTGGA  GAAATAATTT  CTCAAAGCTT  CCCATGTTTG  GCAAAAAAAC  ATTAACTTGC
186301   ATACATATTT  TAGGAGCTCA  ATGAATTCCA  AGTAGGATAC  ACTCAAAGAG  ATCCATACCT
186361   AGACACATCA  TAATCAGATT  ATCAAAGAT  GAAGAAGATG  AATCTTGAGA  GCAGAAAGAA
186421   AGGAACAATT  CATCACATAC  AAATAGTACT  CAAAAGATGT  CTGGAGTAGG  TATACTAATA
186481   TCAGACAAAA  TAAACTTTAA  GATAAGCATT  GTTATAATAA  ATAAAGAAAG  GTATTTTGTA
186541   ATGATAAAAG  TGTCAATTCA  TCAAGAAAAC  ATAACATTAT  AAACATACAT  GCACCTAACA
186601   ACAGAGCCCT  AATATTCATG  AAACAAAACT  GACAGAATTG  AAGGGAGAAA  TAGAAAATTC
186661   GACAATAATA  GTTGGAGACA  TCAATACCTC  ACTAGTTAGA  CAAGATCAAC  AAAAAAATAG
186721   AAGACTTAAC  ACTTGAAAAC  ACCTAACCTG  ACCCTAACAT  AAATCTATAG  GTCACTACAC
186781   CCCAAAACAG  CAGAATAAAC  ATCCTTCTGA  AGCTCACATG  AAACATTTTT  CAGGATAGAC
186841   TGTATATTAC  TTCATGAAAT  AAGTCTCAAT  AAATGTAAAA  GGACTATAAT  AATAGAGTAT
186901   ATATTCTCTG  ACCAAAGTGG  AATGAAGATA  GAAATCAATA  ACTAGGCTGG  GCGTGATGGC
186961   TCACGCCTGT  AATCCCAGCA  CTTTGGGAGG  CCAAGGCGGA  CAGATCACGA  GGTCAGGAGT
187021   TTGAGACCAG  CCTGACCAAC  ATGGTGAAAC  CCTGTCTCTA  CTAACAAAAT  ACAAAAATTA
187081   GCCAGGCCTG  GTGGCATCTG  CCTGTAGTCC  CAGCTACTCG  GGACACTGAG  GCAGGAGAAT
187141   CACTTGAACC  CAGGAGGCAG  AGATTGCAGT  GAGCTGAGAT  CGCGCCACTG  CATTCCAGCC
187201   TGGGAGACAG  AGCGAGACTC  CATCTCAAAA  TTAAAAAAAA  AAAGAAACT  AGAAAAATAA
187261   GAACAAATCA  AACCCAAAGC  AAGCAAGAGG  AAAATGAAAA  ATTTCAAAGC  AGCCAAGAAC
187321   AAAAGGCACA  TTATGTACAG  AAGAACAAGT  GTATAGATCA  CATATTTCTC  ATAGACACAA
187381   TATAAGCAAA  AAGACAGTGG  AGCAAAATTT  TTTAGATTAA  TGAAAGACCT  ACAATTCTGT
187441   ACCAAGCAAA  AAAACTCCCC  CCAAATGAGG  GTGAAATAAG  ACAATTTAAT  ACAGAGAAAA
187501   GAGGAAGGAA  TTTATCTAGT  CATATGTGAG  AGTTTTATGA  TACATTTTGT  ACTGTATATG
187561   TGGATGTTTT  CTATTTCATT  TAAAAAATCA  ACCGTGCAAT  TAAATGGTAG  ATTGTCTTGC
187621   TTCTTTTTGA  TTGACACAGT  CATTAACTAA  AATATTGTAG  TATTTTTTTA  TCTCCCTGCC
187681   TAAAGGCAAT  AAACATCTAA  TCAGCAGACT  AGAACAATAA  AAAATATTTT  TTAAAAGTCC
187741   TTTAGGCAGA  ATGATAAAAG  TCCCTTAGGC  ATATTGAAAT  TCCTATTTAT  ACAAAGGAAT
187801   AAACAGTACT  AGAAATTGTA  ACTATGTGAG  TAAACAGATA  ATATTTTTTC  TCCATAAAAT
187861   GTGGTTGACT  ATTTTCACAA  AAATAGTTAA  CAATGTAATG  TGTGATTTAT  AGCATTTAAA
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187921   AGTAAAACAG GCCGGGCACA AAGGTTCGTG CCTGTAATCC CAGCACTTTT GGAGGCCGAG
187981   GCGTGCAGAT CACTTGAGGA CAGGAGTTCA AGACCAGCCT GGCTAACATG GCAAAACCCC
188041   ATCTCTACTA AAAATACAAA AATTAACCAG GCGTGGTGGT GCACGCCTGT AATCCCAGCT
188101   ACTCTGGAGG CTGAGGCACA AGAATCACTT GAATCCAGGA GGTGGAGGTT GCAGTGAGGC
188161   AAAATTATAC CACTGTGCTC CAGCCTAGGC AACAGAGCTA GACTCTGTCA CACACACACA
188221   CACACACAAA AGAAAAGTGT ATGACAACAA CAGTGCAAAA GAAGCGGAAA TGAAAATAAT
188281   GTTATTTTAT ATAAGTGGTA TACTTTTAGA TGAACTACGA TAAATTAATG ATGTATACTA
188341   TAAACTCTAA GGCAACCACT GAAATAATGA AACGAAGAAT TATGGCTAAC AAGCCACAAA
188401   AAGAAATAAA ATAGAATGAG AAAAAATATT TAAGTTGTTC AACAGATGGG AAAAAAAAGA
188461   GGAAAAAGAG AACAAAGAAC AGATGGGACA AATGGGAAAG TAATAGCAAG ATGATAGACT
188521   TAACTCTACC CATATAGATT ATCACACTTA AGGTAAATGA TCTAAATACT CTAATACAAA
188581   AGCAGAGGTT GTCAGATTGA ATTAAAAAAA CAGACAACAA CAAAAAAAAG CAAAAAAAGA
188641   GCCACAACAT GCTGCCTACA AAAAATTCAC TTTAATATAA AGACACAAAT AGTCTAGAAC
188701   ACCATCACTT TTAACCTTAT TTACTCAAAC CTCCTAACTG ATCCCTATTT ATTTATTTAT
188761   TTATTTATTT ATTTATTTAT TTATTTTTGA GACAGAGTCT GACTCTGTTG CCCAGGCTGG
188821   AGTGCAGTGG CACCATCTAG GCTCACTGCA GCCTCTACCT CTCGGGTTCA AGCGATTCTC
188881   CTGCCTCAGG CCTCCCAAGT AGCTGGGACT ATAGCACATG CCACCATGCC CAGCTAATTA
188941   TTATATTTTT AGTAGAGACG GGGTTTTGCC ATGTAGGCCA GGTTGGTCTC AAACGCCTGA
189001   CCTCAGCCTC CCAAAGTGCT GGGATTACAG GCGTGAGCCA CAGCACCCAG CTCCTCTTCA
189061   TTTATTCTTG CTACGCTTCC TCCAATCCAT TTTGTGCATT TGATGATTTT GCCAGTAACT
189121   TCTTTATTTT TCTGGTAAAA TTACTTATGG GTCACTGAGG ACTGGGATGT TCTTTCTTCT
189181   AGAGGGGGGT TGTGTCTGCT TTTGCCAGGA AGCTGGGGTA CCACCAGTCA AGTATTACTT
189241   TAAACTCAAT TCATGAATTG AGACTTTTTT TTTTTTTTTT TTTTTACGC AGAGTCCTAC
189301   TCTGTCACCC AGGCTGGAGT GCAGCGGTGT GAACATGGCT CACTGCAGCC TCAACCTACT
189361   GAGCTCAAGC AATCCTTCTG CCTCACCATT CTGTATAGCT AGGACTACAG GTGTGTGCCA
189421   CCATGCCTGA CTAATTTTTT AAATGTTTTT TTTAGAGATG GGGCTCACTT TGTTGCCCAG
189481   GCCGGTCTCG AGCTCCTGGG CTCAAGTGAT CCTCCCACCT TGGTCTCCCA AAGTGCTGGG
189541   GTTACAGGCA TGAGCCTCTG TGGCTAGCCA AGACTTTTA TTTTTTAGCC TAAATGTGTA
189601   TAAAAGTTGG CTTGTGGTTA CAACTTATCA GGATTGATGA TCTCTCTCTC TCTCTCTCTC
189661   TCTGTCTCTC CCCACCTCTC TCACATCCCT TGCTCTGCTG AGAAGCAGAG CAAACATTCT
189721   AGCAGTTTCC AGAGAGTAGG ATGGGATTAC TTCTAGTTTA CTTTTATCAT CCTTTGGGAT
189781   CGCAGTATTA CTGGGAGAAC ACAAGTATCT CTTATTAGAC ATACCACCTT TGTAGAATCT
189841   GGACTTTCAT TTTAGACTTT ATTTGTTTTC TACTATAAGC AATTTAAGTT ACAGATCTCT
189901   CTACACACTG TTTAAGTTGC ATCCCATGAA TTTTGATGTG CTTTATTGTC ATTATTATAT
189961   AGTACAATGT ATTTTGTAAT TTTTTGTGAT TTGTTTGGAG AGATTGATTA ATTAGAATGA
190021   TGTTTAATTT CCAAATATGT GTGTTTTTTT CCTACATTTC TTATTTTTAT TGATTTCAAA
190081   TTTATTTCTA CTGTAGTCAG ATTTAATAAT TCATTTATTT TTATTATTTT CATTTTTTTA
190141   GAGACAGGGC CTTTCTGTGT TGCCCAGGTT GTCCCAAAC TCCTAGTCCC AAGCAGTTCT
190201   CCTGCCTCAG CCACCCAAAG TGCTGGGATT ATAGGCACGA GCCACCCGTG CACAACCAAC
190261   AATTCATTTA AAAAGTGGGC AAGTGAACTG AACAGACATT TCTCAAAGA AGGCATACAA
190321   TTGGCCAACA AATATATGAA AGAATGCTCA ACATCACTGT ATTAGTCTGT TTTCATGCTG
190381   CTAATAAAGA CTTAACCTGA GACTGGGGAA TTTACAAGAG AAAGAGGTTT AATGGACTTA
190441   CAGTTCCACA TGGCTGGAGA GATCTCACAA TCATGGTGGA AGGCAAGGAG GAGCAAGTCA
190501   CATCTTACAT GGATGGCAGC AGGCAAAGAG AGAGCTTGTG CAGGGAAACT CCCGTTTTTA
190561   AAACCATCAG ATCTCGTGAG ACTCATTCAC TATCATAAGA ACAGCATAGG AAAGACCCGG
190621   CCCATAATTC AGTCACCTCC CACTGGGTTC CTCCCAGGAC ACATGGGAAT TGTGGGAGTT
190681   ACAATTCAAG ATGAGATTTG GGTAGGGACA CAGCCAAACC ATATAAATAA CTAATCATCA
190741   GGGAAATGCA AATCAAAACC ACAATAAGGT ATCATCTCAC CCCAGTTAGA ATGGCTATTG
190801   TCAAAAAAAC AAAAAATAAC AAATGCTGGT GAGGATGTAC AGAAGAGGGG ACTCTTATAT
190861   CCTACTGGTG GAAATGTCAA TTAGCATAGC CATTATGCAA AATAGTATGG AAGTGAGGTA
190921   GGTTACATAG GGTGGTCACA GCCTCCCTTG AAAGGAAACA AGAAACTTGT CAAATTGATG
190981   GAGAGAACAA ATCTCTTGAC ATTACACAAA CTGCATCTGG GGCTAGTGGT TAGAATATCC
191041   TCAGTCAAGG AGGTAGAAGA GCAGGAGGGA AAATCCCTAA GTTCGTGCAA GTGCAGAAAC
191101   CCACAAGCTG TGTTCTCAGG TTGACATATA CTCATTTTAA TAGTAAGAAA CACACCCTTG
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191161    GGTAGAGAAT TAAAATGCTA ATAATACATG TGATGTATGT ACTAGCGTGT ATGGCAATAT
191221    TGCATGCACA TTCAAGAGAC CACCCAAAAC ATATTTAACA ACAATGCCCA TTCCCACCCC
191281    CTCATGGATA ATCACGTAGG ACTCCCATAA CGGGAGTTTC TTCAGTGTCA ATTGGTGCTG
191341    AAGTAGCCGA CCCTGACTCT GCTATCAGCG TGTACTTTCA CCTTGCAATA AACTCCTTTG
191401    CCTACTTTTA CTTTGGACTG GCTTTCAAAT TCTTTTGTGC AGGGAATTCA AGAATCTGAA
191461    CCAGCCCACT GACAACAGAG GTTTCTCAGA AACCTAAAAA TAGATCTACC AGATGAGGCT
191521    GAAAATCTGC TACTGGCTAT TTATCCAAAG GGAAGGAAAT CAGTATACAA AGAGACACCT
191581    ACATCCCCAT GTTTATTGCG TCACTCTTCA CAAGAGCTGA TATATAGAGT CAACCCTAAA
191641    TGTTCATTAA CAGACAAATG GATAGAAAAT GTGGCATATA TACACAATGA AATACTATTT
191701    GGCCATGAGA AGAATGCAAT CTTGTCATTT GTGGCAACGT AGATGAAACT GGAGAACATT
191761    ATGTTAAGTA AGATAAGCTA GGATTGGAAA GATAAATACT ACATGTTATC ACTCATATGT
191821    GAAAGTAGAG AAAAATTTTT AGCTCATGGA TTTAGAGAAC AGAACTGTGG GTACCGGAAG
191881    CTGGGAAGGG TAGCAAGGAG GGGAGGATAG GGAGAGGTTG GTTAATGGTG ACAAAATTAC
191941    AGCTAGATTG TAGAAATGAG TTCCGGTGTT CTGCACCATT GTAGGGTGCA TATGGTTAAC
192001    TCTCATTTAT TGTATATTTT CAAAAAGCTA GAAAGAATT TTGAATACTC ACAACAAAT
192061    AAATGATAAA TGTTTAAGGT GATGGATATA CTAATTACTC TGATTTGATT ATTACACATT
192121    GTGTACACAT ATAAAAATAT CACTCTTTAT CCCGTATATA TGTACAGTTA TTATATGTCA
192181    ACTAAAAATA AAAGAAAAAA AGAATATGAT CTATCATGAT GTATATATCA TGTGTACTTG
192241    AGCAAAATGT GCATGCAGAT ATTGTGTATA ATGTTCTATA AATCAATTAG CTCAAGATAA
192301    TAGATAGGAT TGTTCAGATC TTCTGTGTCT TTACTGATAT TTTGTCTAGT TATTGCATCA
192361    TTACCAAAAA AAGGGTGTTA AACTCTCCAA ATGTGATTGT AGAATTGTCT ATTTTGTCTT
192421    TTCTTTTCCA TTTTTACTTT ATGTATTTTG AAACTCTGTT ATGACATTTT GCTATGTATT
192481    TTAAAACTTC GTTATGTATT TTGAAACTCT GTTGTTAGAA TCATACATTT ATGATTATTA
192541    TGTTTTCTTG ATGAAATGAC CCTTTTCTAT TGTCGTTGTT TTTGTTTTTT CTGAAATGGA
192601    GTCTCACTCT GTTGCCCAGG CTGGAGTACA GTGGCACAAT CTTGGTTCAC TGCAACCTCC
192661    ACCTCCTGGG TTCAAGCGAG TCTCCTGACT CAGCCTCCAA GTAGCTGGGA TTACAGGCAT
192721    GTGCCAGCAT GCCAAACTAA TTTTGTATTT TTATTAGAGA CAGAGTTTCA CCACGTTGGC
192781    CAGGCTGGTC TCGAACCTCT GACCTCAGGT GATCCGCCCA CCTCGGCATT TTTATTTTAT
192841    TTTATTTTTT TGAGACAGAG TCTCACTCTG TCACCCAGGG TAGAATGCGG TGGTGTGATC
192901    TTGGCTCACT GCAACCTCCG CCTCCTGGGT TCAAGCAATT CCCATGCCTC AGCCTCCCGA
192961    GTAGCTGGGA TTACAGGCAC ATGCCACCAT GACTGGCTAA TTTTTGTATT TTTAGTAGAG
193021    ATGGGGTTTT TCTATGTTGG CCAGGCTGGC AACTGACTCC TTTAACAATA CAAAATATCA
193081    CTCTGTCTCT GGTAACACTC TCTGTCTTAA ACTCTATTTT AGCTGTTATT ATTATAGCCA
193141    TTTTAGTCTT TTTATGCTTT CTGTTTGCAT AGTGTATATA TTTTAATATG TTTATTCTCA
193201    AGTTATCTGT GTTTTTATAT TTAAGATGTT TCTCTTCTAG CCAACGTGTT TGGTTCTTGC
193261    ATTTTTAAGT CGATTCTAAC AATCTTTGCC TTTCAATTGA AATATTTACA CCATTAACAT
193321    CTAACATTAA CATTTATTTT TCTTTCCACA GTACACTGGC TAGCATCTCC CATATAATAT
193381    TGAACATAAA GTGTGATAAC TGACATCCTT ATTTCATTCC TACTCTGAGT GGAAAGGGCA
193441    GGGGTGGAGA AAGCATTCAA CAATTTGCCA TAATTATAAT TCTTTTTGTT ACACTGTTTT
193501    CTTCTGCATT AAAAAATATC ATTACATTTT GCATGAATTA TTAGGAGAAA ATATTTTCCA
193561    ATTTTCCTGG AAAATGCCAT AACCACGTCT CTCAATTTTG TTTCCATCTT TCTTCCACAT
193621    TTTACATAAC CTACATAAGA GACACATTAT CAAGTATATT TTACATGGCT TCTCAGTGTC
193681    TTCTCTGTCT GCTAACAGGT TTACCAAGAG ATGGCACTCT TGTATTTCTG GTGGCTATGT
193741    CCATATCGTT TTGCCTTTAA GACAGCGTAA CTACTTCTTT CACCAGTATT AAAGACATGT
193801    ACATTTGATC TGGTTCTTGT GGATGATTTT AAATGACTCA AGCTAATAAT CCTAATTTTA
193861    CCTAAACACT CCATTATTTT AAAATGTATT CCTTTATGCC CACAATAAAC ATTTATTGAC
193921    ATTAGGCTGG ACATTAGGCT TCTCTATGGC AGACATTAGG CTGGACCCTA GCCATATATC
193981    TATTGAGGGA AAAAAAATTA TTTTCTATAT AAGTTCCAG AAAGCCAAGA TGTGTTTTAA
194041    AAACAAAACA AAACATTACA TTCTAAATGC TGTAACAAGA TAAGAAAAAG TGTTGAGGCT
194101    GAGAGAAGAA CAAAGCAGCA AGCAACTCCT GGAAGGACCA CTGCTGCAGA GGTAATAACT
194161    GGTGAACCAT GTTTTGGAGA AGGAAAAGGT CACCAAGAGA AGGAGGGGGT CCAGGGTGTT
194221    CAGAAAGATT GCATGCATAA AGATCAAGGG TAATAAAAAA AATTCCGTAT TATGTAAATG
194281    TGAAGTTCCA GGACCATGAG CTTGGAGAGC ATGAAGTACA GGAGGAGGGT TGGTTTCAAA
194341    TAAATCTGGG AATGAAACAG TGAAGCCTCT GGCAGAACTC ACATCTCTTT CCTCCCCTCT
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194401  TCCTTGCACA TTCCCTTTAT GGAGTAATTG CAGGGATGGG AAAAGTTCAA AACCACCACT
194461  GAGCCTAGGA AGTGCTAGGG TAAAGTGGAG AATGAACCTG CGTGATTTGC TCATCCTAAA
194521  CTAGGTTCTT CTAGGAGAGC CCTTCCCCAT AAAATCTGCC CTCCTCGAAG GGGCCCAGAC
194581  AGCCTAAGCT CACCTCCCAA AGACCCCTTA CTTGCTGACT GAATCTGATT CCACCCAGAC
194641  ATGGCCTAAA ACCCTTCCAT AACTCTATAG CCAAATTCAA TTTTAGACAG GCCTCATACC
194701  AACCTTTCTT CCTCTAAGTC TGCCACCCTA GGCAATTCTC AACATTCTCT ACACACTTTG
194761  GGGCCATAGA CGTGCTACCA AGTCTCCAGA CCTAGACCTG ATGGAGCAGT GCTGTAATGA
194821  GACGACCACT GGCCTTTGAA CCAGACCCTT CTCTGTGGCT CCTATGCATC TCCAACCTGT
194881  TTTGAGCACT GCTGCCAAGA CATCTTTGGC ACTTTGTTGT GAAGTTTTAA AACTGAACTA
194941  ATCTACAAAA CACCTAACCT TTAAAAATTC ATTGTCATTT CATATCATGA AAGATAAAGA
195001  AAGGCCAGGA AACTGTTCCA GGTTAATAGA GACTAAAGAG ATAGCAACCA AATGCAATTT
195061  GTGATCCTGG ATTGAGGGGA AAAAGTGTTG TCAGAGACAT GATTGGGACA GCTGGTAAAA
195121  TTTGAATTTG AATTTAAAGA TAAAGTATTG AGTAATATAG GAAGATGATT ATCTGCAACT
195181  TTCAAATGTT TCAGTAAGTA TATATATATA TAAAGAGATA TAAAGACATA TAAATAAATA
195241  GATGGATAGG TAGAGAAAAA GCAAATGTAT AATATTAACA ATCTAGGTAA AAAGTATATG
195301  AGTGTTCTTT GTACTGTTTT TCTGATTTTT CTATATGTTT GAAATCATTT TAAAATAAGA
195361  AGGTTTTTGG GGTTTTTTTG TTTGTTTTTT GTTTTTAGAG ACAGCATCTT ATTCTGTCAC
195421  CCAGGCTGTA GCTCAGTGGC CCAATCATTG CTCACTGCAG CCTCAACTTC CTGGGCTCCA
195481  GTAATTCCCC CTACCTCAGG CTCATGAGTA GCTGGTACTT CAGGTGTGCA CCACTGCACT
195541  CAGCTAATTT TTATTTTTTA AATTTTTGTA GAGATGGCAT GTTGCTATGT CACCCAGGCT
195601  AGTCTCAAAC TCCTGCCCCC AAGTGATCCT CCCACTTTGG CCTCCCAAAG TGCTAGAATT
195661  ATAGGCATGA GCCACTGCAC CCAGCCCCAA ATAAAAAGT ATTTTATTTT AATTAACTAA
195721  TTAATTTTGA GTCAGAGTTT CACCCTTGTC ACCCAGGCTG GAGTGCAATG GCATGATGTT
195781  GGCTCACTGC AAACTCTGCC TCCTGTGTTT AAGCGATTCT CTTGCCTCAG ACTCCTGAGT
195841  AGCTGAGATT ACAGGTGCCT GCCACCATGC CCAGCTAATT TTTATATTTT TAGTAGAGAC
195901  GGGGTTTCAG CATGTTGGTC AAGCTTGTCT CAAACTCCTG ACCTCAGGTG ATCCACCCAC
195961  CTCGGCCTCC GAAAGTGTTG ATGAGCCACC ACACCCGGTC TAAAAAGTAT TTTAAAACCA
196021  CAGTCCCACT CTACCTTGTC CTACACTACC AGGGGCTAGG ATCACCCCAT GTCTTCTAGG
196081  CTATGAGATA GAGGAATCCA AGGAAGAAGA TAAGCTACTT GGTTCCTCTA TAGGGTCTTG
196141  TGTGTGCTCT CATGTGCTCT CTCTCTCTCT CTCTCTCTCA CACACACACA CACACACACA
196201  CACACACACA CACACATG AATACCAGAG CTATCACTTT CCCAGTCTAG TACTCATCTC
196261  ATCCCAAGGG TTTTGTGTTG TAGTGGTTTG CTCATTTGTT TGTTTTGTTT GTTTGCTTGG
196321  ATTATTCTTT TTCTCTTTTT GCAGCTGAAG GGAGAATTTC CAGGCCAGCC CTTTGGCCAT
196381  TAGAGTTACA GTGCCTCTAT TCAGGCTTCA TAGAGAGACC TGGGATTCAG TAGTGGGGGG
196441  CTTTTATCCA GTTCAAAATA ATGCATTCTC ACCAAGATGT ACTTTGAAAT AAAACAATAC
196501  TAAAACACAA AATTTTATTT ATGCTGAACA TTGAATCACT TTTTTCTGTA TTTTGTGTAG
196561  AAAGTTATAC ACACACAAAC ACATTTGCTC CTGCTTTGTT TATTGGCCCA GGGGTATGTT
196621  TGGTAATACT TCATCAGGCA TGAGTAGTAC GTCTTGGAAG GTGTGGTCTA AAGCCTAGAC
196681  TCCTATCTGC TTCCTTCAGC ATTCTCCAGT GTATCTGTCA TCTGTCTACC TTAGGATGGG
196741  GTCTCCAGAA CTTCCATTCA CATTTAGAAG AGGGCAGCGG CTTTCTATGG AAAATATGAA
196801  CTCTCATTCA TCTCTATTCC TTCTTCTAGC TATGGTCCAG CTCAGCTGTT TGGAATAAAG
196861  TATCTATATG AAGTCTGCGA ATGGTTCTCA GACTGGTTGA ACATTAGAAT CACCTGAGTA
196921  CCTTCTAAAA TTCTTATTAC CCAGGGCATA TCTCAGAATG AGTACCACAG GGTAGGGATA
196981  GGATTAGGGA TCATGATCTC TGGAGTCTGG TTTAGGCACT AGTGCTGTTT AAAACTACGT
197041  TCATGAGGTG GAGGTTGCAG TGAGCCGAGA TGGCGCCACT GCACTCCAAC CTGGGCGACA
197101  GAGTGAGAGT CTGTCTCAAC AACACAAAAC AAAAAAAACC AACTACCCTT GTGATTTGAA
197161  TGTCCATCCA AAATTGAGAA CCATTAGGTA AGGCCAAGCT GTATAATTAA AGAGCAGTTT
197221  TCATTTGTCT GGTGTGGTGG CAGCTTTTTG ATAAGGGAAG TATTGTTGCC ATCCACATAC
197281  CTGAGCCTCA CTCCTGAGAA CACTGGTGTG TATGTTGCTA AAAATTCCCCA GGTGATTCTG
197341  AGGTTCCTTC CTGGATAAAA ACCACTGACC CTGGGAATGT ACCCACTGCC AATCTCCTGC
197401  GTAAACCTTG GATACTGGGA AGCCTACAGT TGAAAATATT GGGCTTGAGA TCCTGAAACA
197461  AATCTTGTAT TTCATTAAGA CTAATATTTG GTACAGTGCA GCAAATCAAG GGAATTTTGG
197521  TGGCTGAGTT CTTTTAGAAC TTTTGCATTG AAATAGGTTC AAGCAGCAAT AAGTTAAAAC
197581  TACAACCTCA GCTAAAGGAT TAAAAGACAC GTGAGCTGGG TAGGATGAGG TCTAAGATTG
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197641    GGTGTGGCGG CTCATACCTG TAATCCCAGC ACTTTGGGAG ACTGAGGTGG GTGGATCACT
197701    TGAGGTCAGG AGTTCAAAAC CAGCCTGGCC AACATGGTGA AAACCCATCT CTACTAAGAA
197761    TACAAAAAAA TTAGCTGGGC GAGGTGCCAG GCACCTGTAA TCCCAGCTAC TGGGGAGGCT
197821    GAGGGAGGAC AATCACTTGA ACTCAGGAGG CAGAGGTTGT AGTGAGCTGA GATCGCACCA
197881    CTGCACTCCA GCCTGGGTGA CAGAGCAAGA CTCCATTTAA AAAAATAATA ATAATAATAA
197941    CAATAATAAT AATTCAGACA TATCCAGGCA TCAAACAGAT ACCTGGGGCA GATGAATAGT
198001    CTTGAGATTC AAGTCACACA TGAAATTTAG GTGGAAAATG ACATTGGAGA AATTTGAGAT
198061    TATGATGAAT GGAAATTTTT CAAAGAGGAA TTTCAGGCTC TGTTCTTGAG GGGATAGATG
198121    GACTTCCAAC AGCAATAACA CAGGATTAAT GAGGACTTGG GATGTTACAT AAATTAGAGA
198181    TGTTAGATGG ATAAAGAGAT AAAAGTACTC TCTCTAAGAA CATGGGACCA GAGATAGGCT
198241    CACTTCTAAC CATCAGATAT AACTAGCAGA CTAAACGGTC TAAAAATAAA AATCATGCCC
198301    CACTCCTGCT TAAGACATTT TAATTACTCT CAGTAACTCT TCAGTTTTTC TACTGTGTTA
198361    TCTTTAACTA CAGGGTTGGT CTGGGTGTGC AACACAAGAA AGCCTGGCAT ATACATGGAT
198421    TCAAGTGTAT GCCATGTACA GGTATTCTTT CATGTACTAT TTCATGTATT CTTTTTCACA
198481    TCTGTTTTTT CCTTCATTGA AGTCAATGGC TGATATTAGA TTCTACTATT CATGTGTACT
198541    AGTTATATAT AATTGTTACA AAACAAATTA GCAAAAACTT AGTGGCTTAA AGCAACACAC
198601    ATTTATTATT ACCTAAGGTC TGTGGATAGA AGTTCTGACA TGGCTTAACT GGGTTCCCTG
198661    CTTCAAGCCT CATGTGGCTG CAATCCAGGT GTTGGCTGAG TCTGAATTCT CATCAGAGGC
198721    TTGATTGTGG AAATTTCCAC TTCCAAGCTC CCTCAGGTTT GTTGAAAAAT TCAGTTCTTT
198781    GCACCGGTAG AAGCTTCTTG GTAGAGGCTG ATTCAACTTC TAGAGGCTGT CTGCAGTTCC
198841    TGTCACCCAG GGTGGAGTGC AGTGGAGCAA TCATAGCTCA CTGCAGCCTT GACCTCCCAG
198901    AATCAATCTG TTCTCCCACC TCAGCATCCT GAGTAGCTGG GACCACAAGT GTGTGCCATC
198961    ACACCTGCCT AAAAAACAAA CAAACGAAAA AAAACCCCCA GAGAACTTTG TAGAGACAAG
199021    CTGGTCTGGA ACTCCTGCGC TCAAGCAATT CTCCTGCCTT AGCCTAAAAG TTCTGGGATT
199081    ATAGGTATAA GCCACCATAC CTGGCATATG GCAAGTCTTG AGCAGGACAA ATACAGATGA
199141    TTTATGTCTG TCTTCCATGG TATTCTAGGT TATTGTTGAG ATGGTCCTCT ATTGTCTTGT
199201    TCCATCTATT GATTAGATAA AACGTTGTTC CTTCTGTTAT TTTTCAACAG TAGCTTTTAT
199261    GTGTCTCTCT TTATCTTAAA ATTCTAACCA AAGAGCTGCT CTTTTCTTGG TGTACTTTAC
199321    CTTTGGTTGA TCCTTCTTAA CCTCTTCTTG CCCTCTGGGG CCTAAGATGA GGGCTGTTAT
199381    CAGATGTGAG TCTATGGGAA AGCAAGCAAG AGGTTCTTCA GCCTCCGTTC AGCCTTAAAT
199441    GTCTAGGTAG AAATCAGTCA TGGCCCTTCC AATGTGGTAC AGACCAGATC ACAGAGACAG
199501    GGGTCTCAGC CAAGGTCTTG TGGCCTAAGC CTTATAGAAA TAATGAGTGT TTACTTACTT
199561    GGAGAACTCC CTTGGAATAT CTTTTTTTGT GAACCTGAGG CAACTTTTGG TGATTTCTTG
199621    ATGTCTTGGG AATCTTGGTC TAGAGCCATT TCAACCTGAT TTCTTTTCAT GTCAGTGGCA
199681    TTTTGTGACC AGATAGTAAA TAAGTTCTAT GATGTTCACT CAGAGAAATA CAATGACTTA
199741    TGATGTGAAG CTTCTGTGGT TCAGCCCTTA CTTCATCTTC ATTCCCTCTT ATCTGCATCT
199801    GTCTCCTGCT TGGGAACAAA AGTCTGGCTT CATTCTATGA CCCCCACGTT GAGTTTCTTA
199861    GTAGCACTTA CTTTTCAATT AGGAGTGTCC TCACTTCTAT CCATCAGACA TAACTAGCCG
199921    ACTAAACAGT CTAAATATAA AAATCATGTC CTACTCCTGC TGAAAACATT TTAATTACTC
199981    CCCATCATTT AATTTTTTCT ACTGGGTTAT CTTTAACTTC AGAGTTGGTC TTGTGTGCAA
200041    CACAAGAAAA CCTGGCATAT ACATGGATTC AAGTGTATGC CACGTGCATG TATTCCTTCA
200101    TGTACTATTT CATGTATTCT TTTTCACATC TGTTTTTTCC TCTAAAATTT ATTTCCTTTT
200161    AAAAATGAAA ATTTTGCATT TGACTAAATT TGTCAAATTT AGTCAAATTT GTTTAAAACC
200221    ATTTTTAAAA TGTTTCCCGA AGTTTTGAGT GAAGTTAGTA CTTCAGAAAA ACTGTTTTGT
200281    ATTTTTCATG TGACCTCAGT GCACTGCTGT GCATTCCAT TTCTGCGTCC ACACACATTT
200341    GTTTTGAGGA AATATAGGAA CGACAAGATA AAGTTCAAGC TCCTGGACAT TGCATAAAAG
200401    ACCGTCATGA CCTGGTCCTG TTGACTTCCC TAGATTTCCC GCTATTTCCT AAGTTGAGAT
200461    TTTTGGTTTG GATGCTTTGT GTTTTCCTAA AATCAAAATA GGTTTTTGCC TTTTATGATT
200521    ATACAGTAAA TAAATGCTAT TTGTGTGAAA CTTTAAACAA TACAAAAAAA ACCTAAGGAA
200581    GAAAGTCAGA TTCATCTAAA AATCCTTGTG GCCAGAATTA ACTACCTTAG TTATTATTTT
200641    CTCTATCTCT CTCTCTCAAT GTATATTTGG TGTAGGTATA GGGGTGTGTG TAGTGTGTGT
200701    GTATGTATAT ATCTGTTTCT ATTCCTGTAT GTGGATGTGC ACAACGCATC CTGCTTTGTA
200761    CACTACAGTA CTAGCATTTT TCTAATGTAA TTCAATATTG TTGAAAACAT TTTAAAAAAG
200821    CTTGTATATA TACACACACA TACACATACA TGCATGTATG TACATATACA CATACAGACA
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200881   AAAATGTATC CTATGTATAT TCACACATGT ATACACACTC ACACGTACAT AGAGTTTTAC
200941   ATCCATAGTT TATAAATGTT GCTTTTTTTT GGTCACCTTT TTGCTAAGTC TTACACTTTT
201001   TTTTTTTTTT TTGAGACGGA GTTTTGTTGT CATTGCCCAG GCTTAGTGCA GTAGCGCGAT
201061   CTCACCTCAC TGCAACCTCG ACCTCCCGGG TTCAAGCGGT TCTCCTGCCT TAGCCTCCTG
201121   AGTAGCTGGT ACTACAGGTG TGCGCCACCA TGCCTGGCTA ATTTTTGTAG TTTTTTTATA
201181   GAGACGAGGT TTCACCATGT TGGCCAAGCT GGTCTGGAAC TCCTGACCTC AAGTGATCTG
201241   CCTGCCTCAG ATTCCCAAAG TGCTGGGATT ACAGATGTGA GCCACTGCAC CCGGCCAAGT
201301   CTTACACATC TTTTTTTTAC CACTAAACTG TTTACCCAAA CCTGATAACC CAAGTCAACA
201361   GCTATTATGG CTCACACAAT CTTATGTAAA CAAAGATACA GATATATAGA ATTTTCTTGA
201421   TTAATATTCA GAAAAAAATG GAGTCCCTTT ATACGTCCTT AGTATCTGCT TTACTCATTT
201481   AAAAATGTAT TACATTATAT GAAAGTATTC AGGTCAAATG TTATAGATGT GATTCATTCT
201541   TTTTAACTGT GTTATTTTTC TGCAATGACT ATGTATCACA AAGTACTCAG TCTTCCACTG
201601   ATGAAAATTT GGGCTATTTC CAGTTTGTCT TCCATTTTTC TTTCTTCCTC TTGGATTTTC
201661   ACTCAATGTG TTTACTAATT TAGGAAGAAT CAATAGTTTT TATGGTATTA CTTCTCCCAT
201721   TCAAGAATAT AGCATATGGT ATAGTATAGT AGAGTACTTA GTTTAATTTA GCCAGATCCT
201781   GTTTTCTGCC CTTTAATAAA ATTCTATCAT TTTCTGCCTT TGAGTCACAT TTTCCTTGTT
201841   CATATAATTC TTAAAAAATG TATAGTTTTC ATTCTAAGGG AACATAAAAA CTTCTTTCCA
201901   TTTCTATTCC TGTCTAGTTA ATTCTACTAT TGGGAAAAGT AACTGTTAAA AAAAATTCTT
201961   ATCTTTCCAG TCAGTTCACC ACATTTCCTT TATACCTTTG TACTTTAATC CCCAGTCATG
202021   TTGAACACTT CTTATTCCTC ACACCAAGCC TCAACGGGTT TGCTCTTTCT GGAAGGTGCT
202081   TCCCCTGTAT TACTGACTTA TTCATACCAC ACATGGAGAC TGGCGCAGCC CTGTTCTGCC
202141   TGGGAAGCCT TCCCCTGATA CCCCTAGTTG GCAGGAGTCT TCATTTGTTC TTTTCTAGTC
202201   ACCTGTGCAA GTTTGTATTG TTCATGTTTA TCATCCTTCA TTCTAGTTGT CTGTCTCTAT
202261   GTGTGGTCTC ATTCAGTGGA CTCTGAACTC TTATGAAGTC ATGTCATGGG TCAGATCTTA
202321   ATAAATTAAT ATTGTCGGAA GCTAATGTCA TGTCTAGAAT ACAGAAAATT TATCAAAAAA
202381   AAATATAGTA TGTTGGCTGG GCGCAGTGGA TCAAGCCCGT AATCCCAGCA CTTTGGGAGG
202441   CCGAGGCAGG AGGATCACAT GAGGTCAGAA ATTCAAGACC AGCCTGGCCA AAATGGTGAA
202501   ACCTCATCTC TACTAAAAAT ACAAAAAGTA GCCAGGCGTG GTGGTGCCCA CCTGTAATCC
202561   CAGCTACTCA GGAGGCTGAA GCGGGAGGAT CACTTGAACC TGGGAGGCAG AGATTGCAAT
202621   GAGCTGAGAT CATGCCACTG CACTCCAGCC TGGGCGACAG TGAGACTCCA ACTCAAAATA
202681   ATAGTAATAA TAATAATAAT AATTGTATGG AATTGAACTG CTCTGATTGG AAATAGCTGT
202741   TTTTAAAAA ATTATTATTT TTAAGTTCC TGGGTACATG TACAGGATGT GCAGGTTTGT
202801   TACATAGGTA AACGTGTGCC ATGGTGATTT GCTGCACCTA TCAACCCATC ACCTAGGTAT
202861   TAAGTACAGC ATGCATTAGC TCTTTTACCT AATGTTCTCC CACACCCCCA CCCCATCCTC
202921   CCCCAACAGG CCCCAGTGAG TGTTGTTCCC CTCCCTGTGT CCACGTGTTC TCATTGTTCA
202981   GCTCCCACTC ATAAGTGAGA ACATGAGGTG TTTGGTTTTC TGTTCCTGCC TTAGCTGTTA
203041   ATGTCAGGCC AGAGAGGCTT AAATTTTTAA GGATCTCTGG ACTTTTCTTC TACATTACTC
203101   TTGATGTTTA TAAATGTTAC AACTTCTTTA ATTTCATTAA ATGTATACCT TATTGAGTTG
203161   ATTTAACTGA GTTAACTTTG TTATATGAAA ATCATGATTG GGAGTGAGGG GGTTAAACCA
203221   GCTACAGAGA TCTTGATTGT TGGTGGTGAA GCAATGCAAG AATTCATTCA TTCAGTAAAC
203281   TAATGTTTAT TAAGCGTGTA CTGTCTTAGT CTGTTCAGAC TGCTGTAACA AAATATCATA
203341   AACTGGGTGA CTTATAAACA ACAAAAAATT TATTTCTTAC AGTTCTGGAG GTGGGAAGTC
203401   TAAGATTAAG GCCCTGGCAA ATTTAGTGTC TGGTGAGGAC AGGTAGCCAT CTTTTTGCTG
203461   AGTCCTAACA TGGCAGAAGG GTTGAATAAA CTTCCTTGGG TTTCTTTTAT AAGGACACTA
203521   ATCCTAGTGA TGAGGTTTCT GCCCTCATGG TATAACTACT GCCCAAAGAC CCCTCCTTCT
203581   AATATTATCA CTTTGTGGGT TAGGATTTCA ACATGAGTTT TGAGAGGATA CAGACATTTG
203641   GATCATAGCA CACACCATAG GACAGACACT GTGCCAAGAA TTGTGGATAT AGTGATTCTC
203701   AAAATGAACA AGATCCCCTC AGAGAGCTTG CAAAATCCAG CTATAAAATT ATGCTTTTTA
203761   AACAAATTAT GCAGTTTGAA AAATCTACTC TGAATCTTAC TTGTGGCATT GAATACTTTC
203821   GGCCACTCTT TCCTTATTAT ATTAAATATT TACTCTTGTT TGGGGGATCC AGTCTCACCT
203881   ACTTTTTCTA CCAGAACTGG TATCAGCTCA TGCTCTGCCT TATGCAAATT AAGAAAATAT
203941   CATACCTTTT GGGTAAATTA AGCCAAGAAA GTTCTCCTTT CTTCTCTTTC TCTCTTTCTT
204001   TCTTTCTCTC TTTCTCTTTC TTTCTTTCTC TCTCTTTCTT TCTTTCTTTC TTTCTTTCTT
204061   TCTTTCTTTC TTTCTTTCTT TCTTTCTTTC TTTTTCTTTC TTTCTTTCTT TCTTTCTTTC
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204121   TTTTTCTTTC TGACAGGGTC TTGCTCTATT GCCTAGGCTG GAGTGCAGTG GTGCAATCTC
204181   AGCTCACTGC AGCCTTGAAC TCCAGGGCTC AAGCAATCCT CCTGAGTAGC TGGGACTATA
204241   GGCATGTGCC ACAACATCAA GCTAATTTTT GCATTTTTTT GTGGAGACGG GATCTCCCTA
204301   TGTTGCTAAG GCTGGTCTTG GATTCCTGGG CTTATGCGAT TCTCCTGCCT CAGCCTCCCA
204361   AAGTCCTGGG ATTACAGGCA TGAGCCACTG CCCCTGGCCA TTATAACTAT TTTCATTGGC
204421   TTATCAGGCA CATGATAACT ATAATAAATC AATAACCAGA ATTTTTAAAT AAAGAAAGGA
204481   AGGAATTGTT TCAACTCTTC CTGCTACCCC TCTATCCCTC AAAAGGGTAG GCTGAATGTT
204541   GTCCTCCAAA GATATCCATG TCCTAATCCC CAGAACCTGT AAATATATTA CCTTATATGA
204601   CAAAAGGGAC TTTACATGTT TAATAAGTTA AGAATTTTGA GATGGGCAGA TTTTCCTGAA
204661   TTTTGCAGAT GGGCCCTAGT GTAATCACAA GGGTCCTTAT AAGAGACAGG CAGAAGAGTC
204721   AGAATAAGAG AAAAATACTT CAAGATGTTA CACTGCTGGC TTTAAGGTGG AGGAAAGGCC
204781   AAGAGCCAAA AAATGCAGTG GTCACTACAA GCTGAAAAGA AAAAGAAATG GATTTTCCCC
204841   TAAAGCCTCT GGAGGGGGCA CAACCTTGCC AATACCTTGA TTTTGGCTCA GTGAAACCCA
204901   TTTTGGACTT CTGACCTTTA GAACTGTAAA TAAATAAATA ATTTTGTGTT GTTTCAAGCC
204961   ATCACAGTTG TGGTAATTTA CTACAACAGC AATAAAATAG AATTAAATAC AGAGATCTGA
205021   GGAGTTGAGT AGGATAAGCC TACTCCAGCA GGTTATTTCG GGAGTATGGT GAGACTCACT
205081   AGGATGGCGG AACTCAATTA AGGAAGTCTG AAGCTGATAA GCCAGAGAGG GAAGGCTCTC
205141   ACTTCATTTT ATAAGGGTTG CGTCACACTA GGAAGATCCA ATAGCAACCA CAGTCTCAAA
205201   ATTAATGATT ACAAATAGGA CACAATTCCA AGAGTCGGGA GCCAAGCAGA AAATGGATTA
205261   GGGAAGACAT GGATGATATG AAACAGGAAG GAGGGGTACA AGGCAGCTTC CTGGGAAGTT
205321   GCCAGGGCAG TCACAGTTCA CATTCATTAG GCTGTGGGCA CCAAATGCAT ATGGAAAATC
205381   TAGCTGACTT AACTGAACTC CTGAAGAGGA ATGAACACCT CATTTATTGA GGAGCTACTA
205441   CCAATTAGAA TATGTATTTC ATTTGTTCAA TAACCCCATG AGTACAGTAA CACAATCCTT
205501   GCTTTACTAA AGCGGAAGCC AATTCAAAGA GGTTCAGTGA CTTGTCCAAG CTCAGGGAAA
205561   ACACTAGGAA GTGAATATGG GTCTGACTCC ATCACTGATT TCAGGAGCCC TGCCCTTTCC
205621   TCCACACCAT GCCCCCTTGC TTTCAGAAAA AAAGGCTTGT TGACTGAATG GTTGTATGCA
205681   CAGTTCAAAG CAGAAACACA CGATGACATC TTTTGAGATA CTCTAACAGT GAGAACTTGA
205741   AAATGAAGTT AAAAATTAAG CGGCAAAACC AAGCCGAGGC TTTCTGAGAA AGTGGGGCCA
205801   AACCTGTTGC CGTCTGACTG CCACGTGGCT CACTATTTAT CCCTGTAAAA ATCTGCAAAA
205861   GTATTTGAAA GGGAAGAAGG GACAGAAAAC TCCCTCCTTT TCCAAGTTAG CCTTATAGTC
205921   TAGGGCTTAA AATACTGGTT TAATGGTGAA GGTAAGTGCT TTTCTTCTTT TTGGGTAGAA
205981   GGATTATTAC TAACTTACCA AAGGTCCATT AAGGGGAGGG AACAGTTTTA GGAGAAGTCA
206041   GAGAAAAGAC ATTAACAGCA ACATAAGGAT CTCCATCTGG TAATATTGCC TAATTCCAAA
206101   ATGAAGAGAC TCTCTGAAAA AGATAACTGA TTCAATGAAG ACCCTAGGGC AAGGCTTGAG
206161   AAGCCACTGG TACCAATGGA CACTGTGGAC AATGGTCATT TCTCCAAGGA CGCTGTGAGT
206221   ATTAACTGTG ATGCTGTGAT TAGTCAGACT GGGATTGGCT GTGGAATGAA ATACTGATCA
206281   GAACTGACAA GATTTGTGTT TGGGACTGTG GCTAACGAGT CTTTTCAGAC TTCTATATGA
206341   ATTTGAAATG GTCTCTCAGG AAAAGGAGAA CATGGCCGGG CCTGGTGGCT CACGCCTGTA
206401   ATCCCAGCAC TTTGGCAGGC TGAGGCGGGC AGATCACTTG AGGTCAGGAG TTTGAGACCA
206461   GCCTGGCCAA CATGGTGAAA CCCTGTCTCC ACTAAAAATA CAAAAATTAG CAGGGCGTAG
206521   CGGCGCGTGC ACCTATGCGC ATGCATAGTG CGCGTGCCAG CTATTCAGAA GGCTGAGGCA
206581   GGAGAATTGC TTGAACCCAG GATGTAGAGG TTGCAGTAGT TGAGATCATA CCACTGCACT
206641   CCAGCCTAGG TGACAGAGTA AGACTCTGTC TCAAAAAAAT AATAATAATA AAAGAAAAGG
206701   AGAACATGAC CAAAGTTATG AATAAGACTG AAGGCAAGAA AATTGTACGC TTGTAGAGAT
206761   CACCTAGCTT GTTGCCCTCA TTGTACAGCT AAGAAAAGGC ACCCAGGGAC ATTGTGGTCA
206821   GCACCAATTT CTCAGAAAGA TAGGCAGATG ATGAGAGGGC CCTCAGTTTT TCTAACACTG
206881   AAGGAATTGC TTCTATGTTT TCTGGTGAAC TCCTCCCCAC TCATCTTGAG GATTCCAGGC
206941   CAGAAGAATC CACTTTAAAA AAGAAACATT TAAAACCAAT TTAACAACCA ATCAAAGGCA
207001   CTTTTATAGA AATACATTTC ATTTGCTGTT GGCCTGTATT TATGGATCTG AGAGGGCTAG
207061   ACTGCCAATA TTGTGACTGT TTATTATTAT TGCTGTTGCT AGTATCTAGA ATATTATACA
207121   ACATATAACA CTTTGCAATT TACGAGGCAT GTCTCATACT TTTGTTTTCA CTCCAAACTG
207181   CCCAGTGAAG TAACATTATC CCAATTCTTC CTATGAAACA GTGAAAGCCC TAAGAGTTTT
207241   TGAAACTTTA CCTGGTTTAC TCAATTTGGG AATGGCAGAG CAGAATTCAG TCCTTGAATA
207301   TCCTCCCACT GCAGGTTCAT GCTCTTTGAT CTAGGTGTAA CATTTACTCT GAGTAAACTA
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207361  GGACTCTGGG CTAACAGAGA TGAAGCAAGA CAGGCTGGAT ATTAGGAGAA TCTAAGAGCA
207421  ATCTAACGAC CATTATAATA AAATCATGAG TTCTAGACTT AAAAAAAGGG AAAAACCTGT
207481  TTTTTTGCTT ATGCGTATAC CATAATATTT ACATTATTTA TTTTTTTCTC AAATTCAACC
207541  TATACGGTGT CAAGTAATTT TTTTTAATAT AACATTTTCC TTTAACTTAA TTTCAATTCA
207601  TTTTTCTGTG TCTACTTACA ACTTTGGCAC TAGAATTCAC AATTTTTTTT TAGAGGTATA
207661  TCTCCTTAAA GGGAAGGGTT CTGACACTGT TACATGTTCT CAATTGTTTG CAAATAGGTT
207721  AATAATTATT CCAGTGTCTC TAAGTACATA TCAACCATGC CAGTGTTCAG CCTCCATAAT
207781  TTTATTAGCT TCTGTGCTTA TTTTGGAAAA ACATTTCCCA TTACCATGAA AGACCTCAGT
207841  TTAGGATGGT TTGGTATGTT AGCCTGATTT CTGCATTCGT CTCATGCAAA GGAAAATAGG
207901  AAACGAAGAA CTGAAATTAC CTATTGATAC AAAATCAAAG TAGCATTTGA AACCATAAAA
207961  CTTAAGTAGG GCTTTTCATC CTTTCTCGTT AGACAGCAAC AGAGAATGGG AAGAAAAACT
208021  AAAGTGATGG GTTTGTGATA CAATTCCAGT AACATAAAGA GCAAGGAGAA GTAGTTTTGT
208081  TGTGTTTATG TTTAATATTC AAAGCTCAAC CTAAAAGTAT TTTTCATTAT CAAACTTCCT
208141  TCTAGAATAA ATGATTAAAA CTTGATTTAA AATATACAAA TTCTCCTTTA TAATACCTCA
208201  AAATGGAGCT ACCCCATTGA GTTTTAAGCT TGTGATTAAA ATATTACGAA AACAAAGGGG
208261  AAGTTGTAAT AGGTAGAACA AGCAGTAGTC TAGGCATTAG GGGATCTGGT GCTGGCTCTG
208321  TGCATCATGT GGTTTCAGGC AACTTTTCAA ATTTTCTACG CAAATTTTCT TATCAATAAA
208381  ATAAACAGTT GGGCCAGAGG ATCTCTGAGT CTCTTTCAGC TTTCAGTGTT TATAAGATTG
208441  GAGAAGTTGG TGGGAAAGCT TTAAGTGGAG TGTAAGTAAT TGCAGCTGCA TGTACAGTTA
208501  AAGAGTTGCC TTCAGCCAAG CCACGGGATC TTGCATAAAA AGTGAAATCA AATAGAAAAT
208561  GGTCCAAACT CTGGGTTTGA CCACAGATGA CTTCAGCTAG GATCTGAGTG TAGAGCAATG
208621  AGCTGAACTC CTGATATCCA GATGTTAGCA AGACTTGGAG GCCTTCTAAG GCAGAGCAAC
208681  AACCAGTATC TGTCCTGGTG CTGACCTGAT CTTACTAGCA ATTGGGCCTC CATTTGGGTC
208741  CATTGTACAA AACAACAACA ACAACAACAA TAAAATCTCC AAACACCCAA AATTCAAAAT
208801  TTAGATGGAG AGATACTATT CCCAGAATTC TAGAGATATT TGGAAAGCAG AAAACTATAC
208861  TTGCCATGCT GATGAAGTCC AATTATTGCT CTTTTAAATA CATTTAGCTA CTTCTGAATA
208921  TAAAATGAGT ATCTACTAAT TATTTACAAA ATCACTTGGT AAATATAGAA AGTCACAAAG
208981  AATGAAGTGA TCATCCTGTT TTGTAACCCA GAAATAGTCA TTACTGGCAC TTGTGTGAAT
209041  CAGTTTCTAT TCCTGTATGT GGATGTGCAC AGCGTATCCT GCTTTGTACA CTAGAGTACT
209101  AGCATTTTTC TAATGTAATT CAATATTGTC GAAAACATTT TAAAATAGCT TCCATCACAA
209161  TAATCTATCA AATTGACTTG CCAGACTCTC ATTATTAGGT TAATTTATCT CTAACATTAT
209221  GCAGTCATGA GTAATACTAC AAAGGATATT TTTGGACACA ATTTTTCATC TATGCCTTTC
209281  TTTATAATCC TTCATCCTAA GGTCACAGAT TATGAATATC TTTAAAGTAC GGACAAGTCT
209341  TTTAAATTTT GTGTGCAAAA ACAGTGCAAA GCCTTGAATG ATAAAATAGA GGTTTGATAT
209401  ATGTGTTTTT TTGTTTGTTT GTTTTGAGAC GGATTCCTGC TCTGTCCCCC AAGCTGTAGT
209461  GCAGTGGCAC GATCTTGGCT CACTGCAACC TTTGCCTCTT GGGTTCAAGC AATTATCCTG
209521  CCTCAGCCTC CTTAGTAGCA GGGTCTACAG GCATGTGCCA CCACACCCGG CTGTTTTTGT
209581  ATTTTTAGTA GAGATGGGGT TTCACCATGT TGGCCAGGAT GATCTCGAAC ACCTGACCTC
209641  AAGTGATCCA CCCACCTCAG TATCCCAAAG TGCTGGGATT ACAGGTGTGA GCCACTGCAC
209701  CCGGCCGATA CATGTGTTTT TAAAGTCACA GAAATTTCAG ATGTCTTGAA GGATTTTAAG
209761  CAATTTAAAA AATAAAGTCA TAGAAGCTTC AATTTAGGAA TGAATGGAAA ATTGATGATA
209821  TTCTTAGGAT ATGGATTTTT CCTAAAAGAA ACAAATGTAT GCATCCCCAA AGATAATTTG
209881  ATTAGTATAC AAATATTAAA TTAAACATGT CCATATTTAG AGCCATGAAT TCTCTTTGCC
209941  TGTCACAATA GCTGGATTTA TTCACAATTG TAGTAATTAG TCCCTGTTCA TTATAATTTT
210001  CTAGGTGATA TGAAGACTTT GTCAGTCCAA GCAAGTGTCC ACATTGTGTG TAGCAAACAT
210061  GAGAATAAAC ATTTTAAACT TTTAAATGTA ATACATATTA GTGTTATGTA ATGTCATCCT
210121  TCATGTTCGA AGGCACATGG AACATTGTTC TGGTGGTACA GAGGGGAGAG AAACACCATC
210181  AGAATGAAAG GAAAGACCGC TCTGGAACCT TCCTCCTTAG CTCTTGAGCT TAGTTTAATT
210241  GTCCTGTCTT ATGGTCTGCT ACAAGCAATA CCACTCTTCA CCTTCGCATG CTTCTCTGTG
210301  GTTTGATAAA GTACATGCAA TTTTTCATTT AATTCTTCCA GCTGCACTAA GAAAGGAGCC
210361  TTATCTTTAT TGAACAGATG AGGAAATGAA TGATTAGAGA ATTTAAATGA CTAGCTCTAG
210421  GTCACACAGC TGGAACTTAC AGCCAGATTT CCTTTTAACA ATCCTGTAAC CAAAAGCATA
210481  CCAGTAGTGC CCCATAAAAT GTAAGTTATA GAGCTGTGTT GGGTCAAAAC TTTTACTGAT
210541  GCTAAGAGGA GGCAACATTA ACAAGGGGAA ATTATTTGTG TATTATGTTT TGGATTATGT
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210601   TCTCTCCATA GATAAAAGAC TGTCGTAGTA AAAGAGATTC AGGGCACAGG GAAACTCCAC
210661   CACAAAGCGT GGTACCATTT CCCACAGAAG CTAAATGGAC GGGAAGCCTG CCACCAGGAA
210721   AGGTAAAGCC ACTGCTCTTG TTTGCAGGCT ATGTTAATAA GCTGAAGCTT ATTCCGACAC
210781   ATTTACACAT CTCTGCATCA CACTGACCCT TCGTAAAGAT ACTCCCAGTG TAACATTGGA
210841   GCCAGCTCCA GCCCCTGATC CTGTTGCTTT TTCCTTAGCC CCATGAAATC ATCTGCGAGA
210901   AATTAAGCCA AATAAGCAAT AAATCCTGGG ATCTAGGGAG TGGAATAAGT TTTGGGAAAG
210961   TCTTTTTTTT TTTTTTTTTG ACTGAGTCTT GCTCTGTCTC ACAGGCTGGA GTGCAGTGGT
211021   GCGATCTCGG CTCACTGCAA CCTCTGCCTC CCGGGTTCAA GTGATTCTCC TGCCTCAGCC
211081   TCCCGAGTAG CTTGGACTAC AGGCACACAC CACCATGCCC AGCTGAATTT TTGTATTTTT
211141   AGTAGAGATG GAGTTTCGCC GTGTTAGCCA GGATGGTCTC GATCTCCTGA CCTCGTGATC
211201   CACCGGCCTC GGCCTCCCAA AGTGCTGGGA TTACAGGCAT GGGCCACCAC GCCTGGCCCG
211261   GGAAAGTCAT TTTAAACCAA CCTATGTATG AATCCCTACT ATAATATTCT CACCAAGCGG
211321   CTGGCTCTTT CTCCTGAGCT TGGAAACCTC CAGTAAAATG GAAATAATTA TTTCCCAGAC
211381   CACCACTCTT ATCTGTGAGC TTTTTTGGCC ATTAAAAATT ATTTCTTCCA TTATATTTTT
211441   ATCTGTGTCT TCACAGGTTT TCTCTTTCTT TCACTTTAGT GCTTTTCTTC AAATAAGCAG
211501   GAAAAATCCA ATCTATCATG CACATGGGAA CCCTTTCAAT ATTGGTCTGT GGTTGTTCCA
211561   TTTTATGGGG ATGCTTTTAA AGAAAAAATT TGTCCTTTCA ATATATTGAA TATCTTCCAG
211621   CACCACATCA CCTGCAAGCT TTGTAAAAAT AGTTCTACAT ATTAATTTTT TTTTTTTTTG
211681   AGATTGAGTC TCATTCTGTC ACCCAGGCTG GAGTACAGTG ACATGATCTT GGCTCATTGC
211741   AACCTCTGCC TCCTGGGTTC AAGTGATTCT CCTGACTCAG CCTCCCGAGT AGCTGGGATT
211801   ACAGGCATGC ATCACCATGC CTGGGTAATT TTTGTATTTT TAGTAGAGAT GGGGTTTCAC
211861   CATGTTGACC AGGCTGGTCT CAAACTCCTG ACCTCAAGTG ATCCACCTGC CTTAGCCTCC
211921   CAAAATGCTG GGACTACAGG CGTGAGCCAC TGCACCCCAC GTAGTTTTTT TTTTTTTTTA
211981   AGTTGAACAT ATGTGAAGGC AGGACCTAGT GACACATAGC AATAACATTT CCAAGTAGAC
212041   ATTACACTAG GGAATTAGTC AAAGTGCTCA TTTAAAGTAC CATCTCTCAA ATGTATTAAA
212101   AGAGAATCCT TGGATGTGCA ATACCTTAAT TCAAAGGCAG CTCGTTATGT ATAAACTCTC
212161   AAGCTTTGTG ATAAACAAAT GTGCATAACA GATGGGACTA TTGACTTACA GCCCAGGGAA
212221   TTTTATTGAC GCTGAGAAGG TTATGTGACT GGCTCTGCCA CTGTCATCCC CATTCACTTC
212281   ATTTTGGAGC AATATGACAT AAATGCCTTA CATGTGGGTT TTCTCTATTT ATCATGTGTT
212341   TCCTATCCCC TTGAAAGATG GCCATATTTG CTTTACTTGG TTATAAGATC CCATATTCGC
212401   TGTCTTGAAG CCAACCAAAT AATTTGACAA AGTGGGTTTG TAGTGCTGGC TATTTTGGTG
212461   AAAAAAAGAC AATGAGACTT CATGTGTCAT CCAAAGTTCT ATCAGATCGA GCTGTGAGAG
212521   AAAGGAAAAG AAAGGGGTCT CAGTCAGGAT GCTCACTGCA TACATCTGTG TTGTTGTCTA
212581   GGTCCAGATT TCTGTTCATT ACGCTATGGG CTGGCTCTTA TCATGCACTT CTCAAACTTC
212641   ACCATGATAA CGCAGCGTGT GAGTCTGAGC ATTGCGATCA TCGCCATGGT GAACACCACT
212701   CAGCAGCAAG GTCTATCTAA TGCCTCCACT GAGGGGCCTG TTGCAGATGC CTTCAATAAC
212761   TCCAGCATAT CCATCAAGGA ATTTGATACA AAGGTAAGTA TGATGGAAAA TAGGGCTCTT
212821   TGTTGAGAGA AAAAACTTTG AAAGGAAGGC ATAGATCTTG ATTCTGTGGA GTATGGAAGT
212881   ATACATTTCC AATGACAAAT TAAAACTGAC TGGAACTATT TTTCTTTGAG ACATTGCTTA
212941   CTTCAATAAT AAAAATAAGA TTTCATTGAG GTTATTATGA TTATAAGGTG GGGGAACTGT
213001   AGAGTTAAAT GTGAAAAATT TAAAAATGGA ACAGTTTATG TGATGTCTTC AATGAAAAAC
213061   TAGGTATTAC CTGGGCACAT TCTTATAGGT TACTCAATCC TATTCAGTTC TCTGCCTGTT
213121   TTATTGTTTC TGAGCAATTT TATATCCCTG TAAATTCTAT ATAACCAATA GAAATGCAAA
213181   CGATTCTTGT CCATAGCTTT GCAAATAAAT TTTGCCAAGA GAAAAATCAG TTAAAACTTT
213241   TCTCCACTCA CCTCCCAGTT GAATTAGCCA ATTTTGCTGT TTGTTTGTTT GTTTGTTTTT
213301   TGAGATAGAG TCTTCCTCTG TCATTCAGGC TGGAGTGCAG TGGCATGATC TCAGCTCACT
213361   GCAGCCTCCG CCTCCCGGGT TCAAGAGATT TTCCTGTCTC AGCCTCCCAA GTAGCTGGGA
213421   GTAAGGGGGC ATGCCACCGC GGCTGGCTAA TTTTTGTATT TTTAGTAGAG ACAGGGTTTC
213481   ACTAGGCTGG TCTCGAACTC CTGACCTCAG GTGATCCACC CGCCTCGGCC TCCCAAAGTG
213541   TTGGGATTAC AGGTGTGAGC CACTGTGCCA GGCTCTGCTG TATATTTAAA GTCTATTTCA
213601   GCATTGCTTC CTGCTTGTGT TATGCGTGAT CTTTGAGTT TTCCTTTGAA CCAGTTATAA
213661   CATCTTACTT ACTTCCTCCA TTAATCAATG AGTTAAATAA AATCTTTGTT GTATGTTTAT
213721   TTTACATTTA TATGAAAACC ATGAATTTAC CCAATTAAAA AAATTATCCT TTAAATTATC
213781   TTGTACTGTA CATTTCCCAT GTCATCCCTA TAATTCATGA TTAATGATTT TATTACATTG
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213841   GACCTAGCTT ATTTACAATG AGTACATAAA TTTATTGTCT CCAGTCTTTC CTCCATTATC
213901   CCGTCTACAT ATCCACACTG AGTAGATTCA CTACTCAGGA ATCTTGGACA CCTTCAAGTT
213961   GCCAAACATG CAGTGTTCAC TGGACATGCT GTGTTCCTTC AGAATTGGG CCTGCTTCTC
214021   AGCACACTCA CATCTGCTAT CAATGACCCA TGGAAAGTTT TTGCCCTGAG CAAGCCAGAG
214081   TCCCTGTTAG TTTCTTCCAA ATGCTACAAG TTCACTTTTG CTATTTTTTC CGATGAGATA
214141   AAATTTCCT TTTTGACTTT CTACAAATCA TAGTCATTTT TCAAGGGATA GTTCAAGTAT
214201   TGCTTCCTTT CTGGGACCTT CCCAAATTAT TATTTTCTCC TCTCAAAGTC TCTGTTTTAT
214261   TTATGTTCAT CCTCAAATCT TGATTCTCAC ATGAATCATA TACCTTGTAT TATTTATAGT
214321   TTTTTTGAGT AGGTAAAATA TTTCATATTT TATATTCTTT GGCTCTCTAC TTTATAGCAT
214381   GATGCCAGAT ATTTAGGGGC CTTACTGCAT TTATTTTTA TTTTATTTTA AAATCTATTT
214441   TATTTTTTAT TTATTTATTT TAAAATCTAT TTATTTTTAG GTAAATATTC AGGTAATATA
214501   ATTTATGTAA TTATTTAGGA ATTTTAGGTA GTTATTTTAA AATAATTCAA ATTATTTATT
214561   GAGTTATATC AGAAGAATGT GATCTTATTC ATTTGTAATA TGTGTTTTAG GAACTCAGTT
214621   CAGCCAGGGC AGACCATAAT TCCCAAACTT GACTTTTCTT TTTAATTAGG CACTGATTTT
214681   GGTTAAGAGT TCAGTAAAGT TTTGTGTGTG TGTTTTAAAA AATTCTTTGA TATAAGAGTC
214741   AAGATGTTAC TCAACTTTTA CTAGAAGCAA AATAGAGGAA GTGCTTTCAC AGATGAAATA
214801   TCTCTCAATG TTTTCTTCCA TTTACTTCTT CCTATTATTC ATCTATATAA TCATTTTCTT
214861   TACCTCTTTT CTTCATTTCT TCTGTTTTTC TCTCCTACTA AGACAAGCAA ATTAGGGGTA
214921   TAATTGGTTA TTTGGGAAGG TAGGAAGAAT ACAGAGAGAA ACAAAAATCA ATATTTTATA
214981   CTAGGGTCTC ACTAACCTCA AGCAACTCTG ACTGTAAAGT AGATTTCAT AATAGGACTT
215041   CTTGACAAAG AGTTTTCCTA TTTTTCCCCC AGGCCTCTGT GTATCAATGG AGCCCAGAAA
215101   CTCAGGGTAT CATCTTTAGC TCCATCAACT ATGGGATAAT ACTGACTCTG ATCCCAAGTG
215161   GATATTTAGC AGGGATATTT GGAGCAAAAA AAATGCTTGG TGCTGGTTTG CTGATCTCTT
215221   CCCTTCTCAC CCTCTTTACA CCACTGGCTG CTGACTTCGG AGTGATTTTG GTCATCATGG
215281   TTCGGACAGT CCAGGGCATG GCCCAGGTAT CCAGATACTT TCTCATTCTT GGTGGGATCC
215341   AGATTTCTGA ATTCTACAAA ATATCAAAGG TCTTAATGAT TTTCATTTCA GGGAATGGCA
215401   TGGACAGGTC AGTTACTAT TTGGGCAAAG TGGGCTCCTC CACTTGAACG AAGCAAGCTC
215461   ACCACCATTG CAGGATCAGG TAAGTGTGCA CAGATGGGTC ATAGCTTTGT CATCTGTTCC
215521   ATCCCACTGT GTCTTATCTT CTATGAATCA AATGGTTTGG GGAAGAGAGA GAAAAAGTAC
215581   TGCTGAAAAA TTCAACAATA TAAGACACTT GCATCACAAA TAGGAAAGAT GCATCTGTGC
215641   AGTAAAGACA TTGAAGCTTA GAAGTAGAAA AAACCATTGT GAGCTAGGTT TCAGCTCAGA
215701   AAAGCCTTAG TAGTCAGAAA AGCCTTAGTA GTCAGAAAAG CCTTGTCGGA AAAAGTTTAA
215761   ACCTTTAAGA ATTGCACACA TGGAAAAAGA TCAAGTAAGC TATATATACA CCATCTTAGC
215821   AATGATTTTG AAGTGAGAAT TAAGGCTACC ACAGCTCCAG GTGGTAAGGA GAGAAATCAG
215881   GCTGGAAGAG TTTGAAGTTT CTGTATTATT CTAAGCTCTT TACTATTCTA TTATGAGCTC
215941   ATTAATTCTC ACAACAACCC TCTCATATAA GTACCATTTT AAATTCTTAT TTTACAGAGA
216001   AGGGAGTTAA GGAAGGTGGA GATTAAGAAA ATTGCCCAAA TACAAATAGC CAGCAGGTGG
216061   TAGGTCTGAG ATTTAAGCCC ATGCAGATTT TAGCCCCAGA GCAGACATTC TCAATCACTA
216121   TGCTAGACTG CCTTTCCATG GTATGTGATC CTACTCAGGC CTCTACAGCT TTATCATTGC
216181   TGTTCTCCCC AGCCTGTCGT GCTGAGAGTA TATACTCGAA GAGCAGAACT AAAATTCCAT
216241   CCAGCTTCTC ACTCCTAGGT CCACTACACA GCTGCATCCT GCAGACTTTT ACCTCAAGCA
216301   ACCCTCCTGC GTTCTTGCTT CCTTCCATCA TAGTTGTAAC CATCTCCTCT ATTTGCAAAT
216361   ACTATCTGCT GATCTCTCTC TTCTAGACTG GTTTCTTTCA ACCTTCTTCC CACCAAAACC
216421   AAGTTAGCTT GCTAAAATAA AGATGGCGCA TTTTTACTCA CCCGCTTGAG AATTTTCAAT
216481   GTGTTCCTTC ATGCTTACAG AGTAAAGCCT GACCTCTTTA TTGCATGAAT ACAAAAGTTC
216541   TTAGCCATCT GGCCCCAACC TTGTTCCACT CAACTCCCCT GTGCAAGCAT GGCTCCAGTG
216601   GCACTGGACA TTGGCTGCTC TCCACATAGA TCTGCACTGC ACTTCCCTCT GGCTCTGCTC
216661   CCGTTAGTTT ATATGCCTGG AAAGTTCTTT GCCCCTGTTC CTTGTGCCAA AATTCCATCT
216721   ATCCTATTGC ATAGCTTATG TAAAAACTTC CTAAACCTTT TTTTTTTTTT TTTTTTTTTT
216781   TTTTTTTTTT TTTTTGAGA CGGTGTCTCA CTCTTCCGCC CAGGCCGGAC TGCAGTAGCG
216841   CTATCTCGGC TCACTGCAAG CTCCGCCTCC CGGGTTCACG CCATTTTCCT GCCTCAGCCT
216901   CCCGAGTAGC TGGGACTACA GGCGCCTGCC ACCATGACCG CTAATTTTT TGTATTTTTA
216961   GTAGAGACGG GGTTTCAAGC CAGGATGGTC TCAATCTCCT GACCTCGTGA TCCGCCCGCC
217021   TCGGCCTCCC AAAGTGCTGG GATTACAGGC GTGAGCCACC GTGCCCGGCC AAAACTTCCT
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217081  AAATCTTATA ATTATTATCA ATTTATCCTC AGATATACTT CCACGTACAT TGTAGTTTTA
217141  TTATATTTAT ATTTTACATC TTTTTTTTCA AATTGCAGTT TGGGACCCAT TAGTGAGTCA
217201  TAAAATCCAT TGAGCGGGTT AAAATCATTA TTTTAAAAAA TGAGTAGAAT AGAATAGAAA
217261  TTGTTGGAGT GCATTGGACA TGGTAAAGTT AAATATCGAT TCATGAAACC ATCGTTTGAG
217321  GCATATGTGT GTGGTTGTAT GTACAAGTGT TTATGCATAT TGGTGTGTGT GTTATGTTAC
217381  CCTGTAAAAT GCATTCTTA CTATAGGTCT CTGTGAAATA TGTGTCTTGT TGTTTTTTAA
217441  TGTAGACTTC CAAAGCCTAC ATGGCATTTC ACTAGTGACA ATCAATTTTA TTCACATTTT
217501  TCTCTCCAAT TGGACCAGAA GCTCTTTGAG GGCAGGGGCT GTATCTTACC GATTTTTGTA
217561  AGTCTTTCAT TTCCTGCCCC TAGCCTCATA TTAGATCATG CAAGAATGCA ACTGTAATCA
217621  CAAGAAAATG CTAATGGGCT GTGATAGCAG AGAGTTACTG TGACAAACTA AGGGATTTAG
217681  ATTTGGTCAC ATTGGTGTTG AGGAGCCATT GAAGAATCAG AGAGTGTGTT ACTATTATTT
217741  GTTAATTTTA ATTATATCAT ATTACTTTAC TGGGGAAAAT CTGTGAGCTA TTTTAGAAAT
217801  AAATACTCTC ATTGCCCAAT AATTCTAAGT CTGCCACCTC ACTGTTGGGA CATTGTTTAG
217861  GGAGGCCACG AAGTCTCAGC CTTTGATATT TTCATAAGTG TTTTTCTCCC TTTTTCCTTT
217921  AGGGTCAGCA TTTGGATCCT TCATCATCCT CTGTGTGGGG GGACTAATCT CACAGGCCTT
217981  GAGCTGGCCT TTTATCTTCT ACATCTTTGG TGAGTCACTT TCTCTTAAAT CCTAATGCCT
218041  CCATTTCCTG AGCATCCATT TTGGCACCTA CACCACCCAC ATTCTTCCTA TATGAAAGAA
218101  AATGTCCTTT ATCAAATGGA AGATGATAAA AAATGTCAAC GGTTGGTATC ATTTTTAATC
218161  TAGTCACACA ACCTGATTAA CACCTTCCTG GTGGTTCTGG GAAGCCACAC GCAAAAGGTA
218221  GAGGAGTTGA CTATTCACAT GGCACCCACC GACTTGTGAT GCAGTCTTGT CCTTCCATAT
218281  CAAGCACCTT CTGCAGAATC TCTACCACCA CATCTGAAGT GCCTGCTATA TGCAGTTAAG
218341  ATGTCAAAGA TAGTGAAGTA CATTTTCAAT GTGTCTTCAT ATTTCATTAT AATTATTATT
218401  TCTGTCCAAG ATGCCTTTCA CCTGTTCTCT ACCAAGTTAA TCTTGCAAAG TTCAATTCAA
218461  ATGTTCCCTT CCCCATGGGC CCTTCCAGGG CTTACCCTGT CAGATTCTGG CATTCTCTCC
218521  TTTATGATAT TTCCTCTCTA GGTTATGTTG GTGTGTAATT ATTTATTTCT CCTTTTCTTT
218581  CCACTAGACT GTGAAATGCT TGAGGCAAGG AATCCATTCT ATGTTTCAT CACTTGGGTG
218641  TCATCATGGT GCCTGATTTT TAGCTTTAAA ATAAAAGAAT CAGTGAATCC AGTAATTAGA
218701  GGGGATTTAA AGAAAACTAG TCCTCAGAAT CTTTTAACAT AGAATGTTCT TCAAATAAGG
218761  AATTCCAATA ATAAGACAAT TTTCTACACT TGATTTTGTT TTTATAGCCA AATGGTGTCA
218821  TTAAATATAG TCCTGGCCTG AATGGCTTTC TCATTAATGA TGCTAATTAT TTTGGTTTGT
218881  ACATGTTAAC CAGGTATTGT ACAAAAATAT TTCTTTTGGG AATCCATAAT GGATGTATGG
218941  CTTGAATACA AATAATACTG TCTCTTGTAA GTGCATTGGA AATTTTTCCC TGCCACATGA
219001  TTTCATGGAA GGTTGTTTCG TGTATGTATG ACTGCAAACC TGACTATTCA GATCTTCCGC
219061  AACAAGACAA CTTATGTGTG CATTAAGAAG TTGCTGCCTA AAATACATAA CACTGTAATC
219121  ATTGGAGACT TTAAAGTAAT TAATCAGCTA TGCAATGCCA CGCTCCTGTT ATCTCCAGAG
219181  GGCTCTGACA TTGACAAATG GTGGCTTTCT ATTTGAGACG TAATATCTAA AAAGCTTTAA
219241  CAGGTTTGTA GAAGGATTGA AAGAAAGAAT GGGAACATTT AGGTCCTTAT GGTAGAATAA
219301  GCATTAATTG ATTAGTGTGT AGAAGGGAGA GGCATGCCAC TTCAGAGGAA ACTTCCTTCC
219361  CCCAGTAAAC AAATCTACCT AAAAACTAAT TTTATCCCTT CTTCCCAGGT AGCACTGGCT
219421  GTGTCTGCTG TCTCCTATGG TTCACAGTGA TTTATGATGA CCCCATGCAT CACCCGTGCA
219481  TAAGTGTTAG GGAAAAGGAG CACATCCTGT CCTCACTGGC TCAACAGGTA CAGTGCACAC
219541  CTTGTACCTG TGGCCCATGC AGAGGTCTCT AGGGCAGGGT GTGGATCTCC TCTGAGAGGC
219601  ACCATCTTGG CTGCTCTAAT ACTCATGCTG ATTAGATCTT TCTTTTCAGC CCAGTTCTCC
219661  TGGACGAGCT GTCCCCATAA AGGCGATGGT CACATGCCTA CCACTTTGGG CCATTTTCCT
219721  GGGTTTTTTC AGCCATTTCT GGTTATGCAC CATCATCCTA ACATACCTAC CAACGTATAT
219781  CAGTACTCTG CTCCATGTTA ACATCAGAGA TGTGAGTTTA CTTCCTATAC TTCTACGAAA
219841  ATGATAATGG TAATAAGGAG AAACAGTTCT GTGTTACCTA TTACATTCTG GCTTTACATA
219901  TAACCATTAA TTTAACCTTC ACAATGACCT TGAGAGAGGC ATTGTTATAA TTCCCTTTTC
219961  ACAGATGTGG AAACAGGACA CTTAGAGGTG AGATAACTTG CCCCAGGTTG CACAATACTA
220021  AGTGATAGAG CTGCTGCAGC ATCCATATTC TTAACCACTA TGCTATACTA CCACACCAGC
220081  TGATTCCAAA GCTTCTTTTA GAAATAATAT TGCTGGGCCA GGCATGGTGG CTCATGCCTG
220141  TAATTCCAGC ACTTTGGGAG GCCGAGGCAG GCAGATCATG AGGTCAGGAA TGCAAGACCA
220201  GCCTGACCAA TATGGTTTAC TAAATATCAT CTACTAAAAA TACAAAAATT AGCCAGGTGT
220261  GGTGGCAGGC ACCTGTAATC CCAGCTATTC AGGAGGCTGA GACAGGAGAA TCGCTTGAAC
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220321  CCAGGAGGTG GAGGTTGCAT TGAGCCAAGA TCATGCCACT GCACTCCAGC CTGGGCGACA
220381  GAGTAAGACT CCGTTTCAAA AACAAAAAAC CCAAGAAATT AATATTGCTT TTATCTGGAG
220441  CCCAGAGTGA TGCAGCTTCT GGCCCTCTTA TCTGAGACAG TGTTCTTTTA GTGTGAAAAA
220501  GGATGCTAAT TTTCCCCCAA ACAACCCACA GTATCATGGG GGTAAGTTAA TGGCTGGTCT
220561  GTGTAACTGA CAAATTTTGG TGCTAACGTA TCTCTATAAC TACTCTGTAT AAACTTCCTT
220621  CCTTCAGAGT GGAGTTCTGT CCTCCCTGCC TTTTATTGCT GCTGCAAGCT GTACAATTTT
220681  AGGAGGTCAG CTGGCAGATT TCCTTTTGTC CAGGAATCTT CTCAGATTGA TCACTGTGCG
220741  AAAGCTCTTT TCATCTCTTG GTAAGGATAA GCGTGTGGGC CCATTTAACC AATCCCTTTT
220801  CTGCACATGG TCTCAGAGGG TTCCCTGACA GCATGTCCTC ATTGCCCAGG GCTCCTCCTT
220861  CCATCAATAT GTGCTGTGGC CCTGCCCTTT GTGGCCTCCA GTTACGTGAT AACCATTATT
220921  TTGCTGATAC TTATTCCTGG GACCAGTAAC CTATGTGACT CAGGGTTTAT CATCAACACC
220981  TTAGATATCG CCCCCAGGTA AGAGCTCTAC CTGTTTTTTC CCCTCCTCCA GACCCCTCCA
221041  GAGGTGTTAG ACCTCAGTGG TCGCCGTGAA ACTCTTTAAT GTTACTGACA TTGCACTAAT
221101  GGCAGAATGA CAAATAACTA CAAATATCTG TCTGTGGCCA TTTTTAGAAC AACAAATGTG
221161  GCATTTTTAG AACAACAATT TCCAATCTTG GCCAGTAATC ATTTTGACAA AAACCTTCCC
221221  AAGCTTCCCT AACAGAGATT GAACTGTGTA TGCTGGGAAA AGGCCCACAC ACAGGTGATT
221281  TGGAAAAGTT TCCATGGTGT TGTTCATATT AGCTACCACA TATATATATA TATATATATA
221341  TATATATATA TATATATATA TATATATATA TACAGTCACA ATAAGCCAGC TCCTGTGCCA
221401  AGACTTGCCA TATATCAACA CATCTAATCC TCACAGTTAT ATTAGGTAGG CCCTATTGTT
221461  ATCCCCATTT TATAAGGGAG AAGGCTGAGG CACAAGGAGG TTAAATGGTG TGACTATGGT
221521  CACATAAAGG CAGAGCCAGG ATTTGGACTG GGGGAGTCTG GCTTTGGAGT CTGTGTCCTG
221581  CCCGTTGCAC AAACTGGCTT CTACACTGAG CAGCCAGGGT AAAGAAACGT GGTTCCCAGA
221641  GAGACTGCAT TGCTCCCTGG TTATTGACTT GGTAGATTGG TAATTTCAGG TTTGGCAAAT
221701  AGACATTGCC CTGAATGTCT TTAGGTGAAT GAAAAACTGC ATTAAGCAAA ATGACTTTGC
221761  CATTAGAGCT GAATTGCATT AAAGTTGAGT TGCTGCAGAA GCTGTAGGTG GCTTTCTATA
221821  TAAAATCATT TATAAAATCA TCTTCCCATA GATATGCAAG TTTCCTCATG GGAATCTCAA
221881  GGGGATTTGG GCTCATCGCA GGAATCATCT CTTCCACTGC CACTGGATTC CTCATCAGTC
221941  AGGTTGGGTC AGTTATTGA ACATCTTCAA GTGGCAGGTA TTGTTTTAGG TGTTGGAGAT
222001  ACACACGGTG CTCTAAAGAT CTGGATGGCA ACACAATTAC TCTATTTACA TGAGCCTCTA
222061  AATCAGACTC TGGTAGGTCA GATTCCCAG AGGAAGAAAA ATATAAGCTT ATTTTCTCAA
222121  GATGAATAGA TGTTAGATTG ATTAAAATGA GCTGTTCCGG TGCAGAAGAC AGCACGTATG
222181  ACTTCCTAGA GGTACATGAG CATGAAACAG TTCTTAGTTA TGACCAGAAT GAAAGACACA
222241  TGTCAAGGAA TAGCAAGAGA CGAAGACAGA GGGGCAAAAG AAGATCATGA AGAATATGTT
222301  CAGACTAATC CAATTTTTAA AAAATCACAA AAGGGAAACA AAGTGTCCTA GGCCAGTTTA
222361  AAGATAATTT AATGTCTGGA AACAGATCGG CTGTGAGACA TTGCAAGGAG GCTTGCTCGG
222421  TGTTTGGAAA TGCAGGCTCA TGAGGAAGAT GAAAAGACAG ACCCAGGCAG GGATGGAAGG
222481  ACTGACTAGA ACCAACTTAC AAAGAGAAGT TTTGTTTTTA CTACATTTCT ATGTGATCAA
222541  GTTCCCAGGT TAATATTTGA CTAAACTGCT AGGAATCCAC TGTGACTATA ATGCTGGAAA
222601  TGACTTAGTA GGGCTTTCTG AGGAGGGTCA CACAGAAGAC CAAAGAGAAC TCATGTTGAA
222661  TTGAGATGGG TTATAGTGAT AGTTGTCAAC AGCCAATACA GAAACAAAAA AAAACAAAAC
222721  AAACAGCAAC AACAACAACA ACAAAAAAAA AAAACAGAGA AGACACAAAC ACAATGCCAC
222781  AATGCCATTT TAGGCATAAT TTTAAATGAG TAATATTATA TGTTGAAATC CAAATTTTCA
222841  GAAAAACATT AGTGTATTTT ATTTTTGTTT AAAGAAATAA CCATCTCAAC TCAGAACCCC
222901  ATGTGCATTT TGGCCATTTT GTTTCCAATA GTTTCATAAA CTTTCTTAAG TAACTACTGC
222961  ACATTGTTCC TTATATTCCT TGTGATCAAC ATTGCAATAC ACAACTGGGA GGGCTACTAG
223021  AACTGGTGTA GAAGGAACTT GTGAGATTGA TCATTTTCTC TGTTTTTTAC ATCTAGGATT
223081  TTGAGTCTGG TTGGAGGAAT GTCTTTTTCC TGTCTGCTGC AGTCAACATG TTTGGCCTGG
223141  TCTTTTACCT CACGTTTGGA CAAGCAGAAC TTCAAGACTG GCCAAAGAG AGGACCCTTA
223201  CCCGCCTCTG AGGACATAAA GTTACAAACT TAAATGTGGT ACTGAGCATG AACTTTTTAA
223261  ACATTTTTTA CTTCTCTCCA TATTCCTGAC CATAGACTCA GCAGTTCTTA ACTCTGGCTG
223321  TGTGTTAGTC TTCCCTGGGG AGCCTTTATA AGACACTGAT ACTTGGGACC CACTCCAGAG
223381  ATTCTGAATG AATTGGTCTG GGGTGGAACC CAGATACTAC TAATTTTTAG ATACTCCTTA
223441  GAGGTTTCTA GCATGCGCCC GGGGTTGACA ACAGCTGGAC AAACTTGAAA AGTCAATTCA
223501  TGTGGCCTTT GAATTTTCCT CATTGGAAAG TACTAAATAA ATAAAAATTC ATGTGAAAAT
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223561   GATCACTGAT AAATATCTTC ATGGTGGGGC AGGTTATTGG ATGCAGAGAA GATCTGCTCG
223621   GAATTGTAGC CATATGTTAC AGATCTCAGC ACCGATCAGA ACTGTAAAGC TATAATCCCC
223681   AGAATTAAAG TTTTTATTAT TTTTTATACA TTGTAAAACA TAGACGTTTA TTTATGTGAT
223741   TAAATTCTAT TAAAATTTAC ATGCTAAAAT AAAATAGACC ATTTTCAAAT TATTTAGATC
223801   CAGATATTTC CATCAGATTA AACAGATATT TATTTATCCT AGCCCAATTG CAAGAGATTA
223861   ATGATGAGAA AATGACCAAT ACAAGATTAA ATAAATGAGG TTAACTTAGA AATCAAGGAC
223921   AGAGAAGATA GAACTGGAAA GCTTGTATTG TGAGAAGAAT GAATGTGAAG GAAGGCAATG
223981   TAGACACTTC CAGAAGGGAT AGCAATATAG TTTAGACCAT ATAATGAAAA TTGGAGAGAG
224041   ATGACAGAGA CACTTTCAAG TGAAATGACA ATTTATATGG GGGAGAAAAA TATTGAAGAC
224101   ATAACAAGAT GAGAAAAGGC ATAGAAATGT ATCACATACA AGGCATAGAA GTGTATCACA
224161   TACAAGAGAA GTTCCTTTTG AGCGTAGAAA AAGATAATTT AACCTTCTTC ATATTTTTCT
224221   TACTTTCCCA AGATACTCAG ATAGGCAGCG TCAACTCTAA CAGGAATTAA TTTGGCTCCT
224281   AACACTTAAG ACATATCCTT TAGTTTGTCT CCTCACACAG AACTGATTCT GGTTTTGCCA
224341   CAACATGTCT AGAGAAGAAG TTCCCACCAT ATTTTAAATC CTATTAAAAA ACTGCTTGGA
224401   CAAGAACCTT GGGCTAATTC AGCAGATGAA GAGAATCTCC TAATGCAAAT CAATGGGTAT
224461   TTTTGAGCAA GTTTTTCAGA AAAACAGAGT GTCAGCCCCT GAGGGTGGTA CTAAGATGAG
224521   AACATTGATT TTGCCTTCAT GATATTGACA ACACAAAGAG GAAAGGGGGT TTGCAGAAAA
224581   CTAAAAGAAG AAGTAGAAGA AAAAGAAAG ACATAGTATA ATAGGTAGTC AAATTATGTA
224641   CAGAAAAAAG AGGAAAAAAA ACCAAAAAAG GGTGGGGGAC AGACAACCCA ACTAAAAAAT
224701   GGGCCAATGA CTTGAACAGG GACTTCATAA AAGAGAAAAT GTAAGTGGCT CCTTAACATA
224761   TAAAAAGATG TTCAACTTCA TTAGTCATTA CAGAAATGAA AATCAAAACT ACAATGAAAT
224821   ACCACTATAA AATTAACTAA TGGATAAAAT GAAAGGAGAT GGAAACAAA ATGTTGCCAG
224881   ACATGTGGAG CAACTGGAAC TTTCATACGT TACGAATGTG AACTTTGGAA AGCTGCTCGG
224941   CAATATCTCC TAAAGCTAAA TGTACAATTC CAGTGACTCA GACATTTTAC TTAGAAATGC
225001   ACATATACAT CCATAAAACA TGTACAACAA TGTTCATAGG AGCACTATCT GTAATAGCCT
225061   GAACAGGAAG TTGTCTGTTA AAAAAGAAT GAGTAAATAA ACCACGGTCT ATTTGTATAG
225121   CAATGAGAAT TAACAGACCC CAATATATAA TAGATGAATG GGTCTCATAA GCACAATATT
225181   GATTAAAGGA AGACAAAACG CACATTCTTT TAAAGGTTTA TAAAATACTT TTTAAAAACA
225241   GCTACAACCA ATCCGTCCTG TTAAAAATCA GTGAGCGATT TCCCTTGTGC AGGGATGGGG
225301   GTTGTGGCTG GATGGATGGT ACTTAAGAAG TGCTCCTGGG GTACTAGAAA TATTTTATTT
225361   CTTGACTTGG ATGTGTGTTT ACTTTGTGAA TATTGTACAT TTATGATTTG TGCACGTTTA
225421   TGAATGTAGA AAATAAAACA GAAAGCAAAT TCAAAGTATC ATCCTTTTGA GAGCTTCTGC
225481   TCTGACTTCG TTTTGACCAA TGGAGCAGTT GGGAAGGGGT CTTGGTCCTT CGGTCCTTTG
225541   CTTTTTTTTT TTTTTTTTTT TTTTAGACAG AGTCTCACTC TGTCGCCCGG GCTGGAGTGC
225601   AGTGGCTCGA TCTTAGCTCA CTGAAAGCTT TGCCTCCCGG GTTCATGCCA TTCTCCTGCC
225661   TCAGCCTCCC CAGTAGCTGG GACTACAGGC ACCTGCCACC ATGCCCGGCT AATTTTTTGT
225721   ATTTTTTAGT AGAGACGGGG TTTCACCATG TTAGCCAGGA TGGTCTCGAT CTCCTGACCT
225781   CGTGATCCGC CCACCTGAGC CTCCCAAAGT GCTGGGATTA CAGGTGTGAG CCACCGCGCC
225841   CGGCCCCTGG TCCTCTGCTT TCATGTTCTT CTTGGTCCTG TTCCTCCTCC TCTTTTGTTG
225901   GAACTTCCAG TATCAGAGCA GGAAGGAAGG CAATGGGTCA ATCGATGCTG TCAGCTTTTG
225961   GATCAAACTG CAAGTTCTCA AACAGCAAAA TTAATGAGCT CAGGCTTTGA AGAAACCATG
226021   ACCCTGAAAG CATCAGTTGC TTCCAATTGC ATCAGTTGCC ACGGGTGATA AGAACAATGA
226081   TGACTCAGAA TGCCTAGGTT TTCCCAGCAG CTTCTCTGAG GTTTTCCCAG CAGCTTCTCT
226141   GATTGATTCC TGACAGATGA CTTCGGTGTG TCAGACTTTC AGGGTATCTT TCCTTATGTG
226201   ATGGTTTGAG GAAGAGTTAC CATTCACATT CCTAATGGCT TCAGAATAGA TGCAATTGTG
226261   AACTGATAGG AAACATTTCT AATTCATCTC CCCTCCCCAT CCCTAAGGA TTGTTTCTAA
226321   CAATAGTCAT GAAAATTAAT TCACTTTTCT CAAATAGTTT ATTGTCATCT ACCTAATGAT
226381   GAGATGACTT ACTTTTTCTC CTTGACTGTT AAATATTATG AATTATATTA ATGTATTTCT
226441   TAATGTTGAG CTTTCCCTTG AATATTCTTT TGATGTACGA CAGAATTTGA TTCACTAATA
226501   GTTTATTTAG GACTTTGGCT GATGTACTGA TATATGAGAT TGGCTCTGTA TGCATACATG
226561   TGTTTTGTGT ATCTTTTTTG TGTCTGGATA TGGAGCTTAT GCTGATTTCA AAAACAAGAA
226621   AGGAGAACTT TCCTTTTTCC CCATTACTCT GAAAAAGATT GACTAGAATG GAATTTTTAT
226681   AATTGCTGTT GTTATTGAA AGCTTGAAAG CATTGGTTTG TAAAAATCAT GCAGGCTGAA
226741   AGCCATTTTG AGGAGACTTT GATAACTTTC TCAATTTCCT TCAGTTACTG GTCTTTTAAG
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226801    GGGTTTTATA TTTTTCTTTG ATCAATTTTG ACCATTTATG TTATCTTGGA GGATCATCTA
226861    TTTTACACAC TATTTAAAGT ATATTTGCAA AAATTCAACT GTTTTATCAG GCTATCTTTT
226921    TAATAATATA TTCATTTTAT CTATATCTGA GGTTTTAGCT TCTTTGTACT TCTGACCCAA
226981    TTGCATGTGT GCTTTCTTTC TCCTTCATTA GACTACTTAG TCATTTACTA ATTTTAAGAA
227041    TAGCTTGTCT TTTATTTATT TACTTATTTA TTTTTGAGAC GGAGTCTCAC TCTGTCACCC
227101    AGGCTGGAGT GCAGTGGCGC GATCTCGGCT CACTGCAACC TCCGCCTCCC GGGTTCAAGT
227161    GATTCTCCTG CCTCAGACTC CCGAGTAGCT GGGATTACAG TCATGCACCA CCATGTCTGG
227221    CTAATTTCTG TATTTTTAAT AGAGATGGGG TTTTGCCATG TTGGCCAAGC TGGTCTCAAA
227281    CTCCTGACCT TAGATGATCT ACCCACCTTG GCCTCCCAAA GTGCTGGGAT TACAGGCATG
227341    AGCCACTGCG CCCAGCCCTG CTTGTCTTTT TATTTTATAT TTGATTAGCT TTATCTTTTA
227401    TCAAGCTTAT GTCCTATTTC CCTTTGCTTT ACTTCATATA AATTTTGTTT TGGATAGTTT
227461    ATTTATTTTT CATTTAATTA TGAAACAGGT TAAAGCTTAG AGGAAAATTG CTCCTCTAAG
227521    TCCACTTTTG TGGGCAGATT ACATTTGCT GTGTTGTGCT CCCAAATTCA TTGTTCTTTT
227581    AATGCTTTAT TTCTCAAGTT AATAACCTAT ATAGTAAAAA AGTGGCTGTT GACTCTCAGC
227641    TTTTTTTTTT TTTTTTTTTT TTTTTTGTA GATACAGGGA TCTTGCTGTG TTGCTCAGGC
227701    TGGTCTGAAA CTCCTGGCTT CAAGGGATCC TCCTGCCTTG GTCTCACAAA ATGCTGGGAT
227761    GACAGACATG AGACACCATG CCCAGCCATG TCTCTCTCCT TATATATAAT AAGAAAACAG
227821    ACACACTGAG GCATCCTATC ATCTCACTCT TGGTTTCACT ACTGTTCTCT GGAAGTTTTG
227881    CTCTGACCTT TTGCAGTTAA TGTATTAATT TTGCATTGAG TAGTTTCCAT AGAAGAATTA
227941    TAGCATTTGC ATTCTGTTGG GTATTATACT TTTCACTGTT ATTTGAACAT AATTTGAGGG
228001    CTGAAACCAA GATGAGGCAA GTGAGGTGCC CAGGAAGCAA TATTTAAGGA GGCATCCTTT
228061    CTTAGGCTCA TGCAAGAACA GAATTGGCAC ATGAGAGTGA GTGCCTCCTT AATTTTGAGT
228121    GCTGGACACT TCTTGCTCAC TTAGCATACC CCTGGACAAT GAAGTGTTTT TTGTTTTGTT
228181    TTTTCATGTC CATCCTTTAT CCTTCTTCAT CTCAAAACAT TTCAATGGAG TATTTTTTTG
228241    GAGCAGTACT TGGATGAGCC TCTGAGTCCC ACAGTAGCTG AGAATTTATT TCATAGTACT
228301    CTTTATGATC ACTGTGGAGC CTTAAAACAT TGTAATATTA ACTTAGCTGG GAACAGAAAT
228361    TTTGTTCCAC AATTTGTCTT ATTCAGAACA GTATTGACTT CCTGCTAGTC TCTTCTGATG
228421    TCCAATATGA GGAAGTCTAG TTAGCCAGCT ACTTTTTGTA GGAGAGCTAT GTTTAGGCTA
228481    GGTGCTATAG GATTCTCTTT ATCCTGGAAT TCCTTCACCA AGATGTGCCA AGGTGTTAAT
228541    CATTTTCTCT TGCTTTTTGG CTGGTGGTCT TAGAGTTTCC TTCGATTTTG TTTTATTTAG
228601    TGATTGTCCT CAATTTGTTT TCTTTACTAA GAATCTCTCT TCTATTTATC TGTATGGTAA
228661    AACCTTGTTG CCCATCTTTC TGGTTTCTGC TGACTTTCAT TTTTGGACCT TTTACTTTGC
228721    TTTCTCCATG GACTTTTTGG TAGTGGAGGC AGGCAAACAC TTTCCAAAGT CTTTCTCAAT
228781    TTCCATCAAT TTCAACTTAT TTCCTAAAAT TGCCTCAGAA TGTGCCTATG TCCACAATAT
228841    CCCTCCTTCC ACTTTAGAAA GGAAAGGCAT CCACACTTTA TTTAGGTGCA ATGCCTGAAG
228901    TGTAAACACT TTCTGGTTGT CAACAAAGGA GTACTTCCAA ATATTGGTTT GGGGATAACC
228961    TGCTAATGAT TAACACATTC ACCTTGGCTC TTGGTTTGCC TGCTCCCTCT TCTTTTATCT
229021    GCTGTGTGTA TTTTTTTTAA TCACTGAGAA TATGCACAGT ATTGTATGTT TTATTATAAG
229081    AGAGGACTGG CCAGAGTGGG AATGTTCTGA ATTCAGAATA ACTGAAGCAG TACAGGATAG
229141    GAACTCATTC TTTCAAATGA AGCTGGCATA TTTTCCCAGA GCACCAAATT TCAATATATA
229201    TTTAAAAAAC TTGATATGAA TGATACAATA AAGTGGTTAG AACTTTTATT AAAATAAACT
229261    TATGTCATGA AATACTTATT CTAATTATAG TCACTCTTCA TCTTATTTCA TCTTATAACA
229321    TGTTTAATGT TTTCTTTTAT TTACAAAACA ATTTATTTTT TGATGAAAAG TTTTAGAAAT
229381    CAAGTTAAAA ATATTCAAAG GAATGCCTAA AGTTTCAAA ATTCTTTTAC ATGTTGTACA
229441    ATCAAAGAG TCTGAAGACC ATTTAGCTAT CCAAATTGTT TATTTTTAAG CAGTATCCCT
229501    TCTAATATTT ACTATTTATA ATCCTTAAAA ATTTGCCTTA GCACAGGAGA ATTGCTTGAA
229561    CCCAGGAGAC GGAGGTTGCA GTGAGCCAAC ACAGTGCCAC TGCCCTCCAG CCTCGGCGAC
229621    AGAGTGAGAC TCTGTCTCAA AAAAAAAAA AAAAAAAAA AAAAAGGCC AAAAACAAAT
229681    AAACAAACAA AAAAATCCGC CTTAACATTA TTTGTTCATT AAAAACTTTC TTTAATACTA
229741    CTAGTTTCCC TTTCCTCTCA GCCCATTGTC ATATTTTGAT TTTTATCACT TGCTTTGTAG
229801    GACATATGAG GTTTTTGTTT TTTTTTTTTT TTGGAGATGC AGTCTCCCTC TGTTGCCCGT
229861    GCTGGAGTGC AATGGCGCAA TCTTGGCTCA CTGCAACCTC TGCCTCCTGG GTTCAAGCAA
229921    TTCTCCTGCC TCAGCCTTCC AAGTAGCTGG GATTACAGGC ACCCACTACC ACGCCTGGCT
229981    AATTTTTGTA TTTCTGGTAG AGACGGGGTT TCACCATGTT GGCCAGGCTG GTCTCGAACT
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230041  CCTGACCTCA AGTGATCCAC AATCCTTGGC CTCCCAAAGT GCTATGATTA CAAGCATGAG
230101  CCACCTGCCC AGCCAGAATA TATGTTCATT TTGAGTCCTT TAACAAAGTC ATAAGAATTT
230161  TAGGAATTCA GTTACTTTCT TGAGAAAATC TCTGAAAAGA TGCCAATAAT TTGTAGCCAA
230221  TTATATTGAT TTCTCTTTTT CATATTGAGA ATTGTTTTTT AAAAAGTTTG TATGTGTGAA
230281  GATTTTTGCA CTGTAGTTAA AGAAACCACC TGTGTGTTGG TTAAGCCATA AGTACATGTA
230341  TTCAAATAAA TTGAGGTGGG GTTACTCTGA GAATCAAAGG AAAACCTGAA GAAACAGGCA
230401  GCCTCAAAAG GTCTTAGCTG TAGCAACTTG CTCCATTGTT GAAATAAATA GGCTTGAACT
230461  TGTATTTTCC CTCTACTCAA CATTTAAGGT CTCAGAAGAT AATATAATTG GTGAAATTTA
230521  AGTAAAGTGC TCACTCTTTT GCTTTAACAA ACCCTAGAGA GCTGGTAGGC AGAGCCTCAA
230581  CAGACCGTTT TAGCTTCCAA AGGGAGTTCA GGACACCATG ATTCACGACC ACAATACATC
230641  ACACATAATT GAGAAAGAT AGTTCCACCA AATAAAGTTG AAATGCTGAC AAGAAGGGGT
230701  AAGAAATCTT GGAAATAGGT TTATATAAAA TTTATTTTTT CCTTTTTTAT TGTTATGGAA
230761  TAGGACCAGT TCTACTTAAG CCACCCATTT GCCAAAATAA AGTGAGAATC GTTTCTTTTG
230821  GGGACTCCTC TTTGTAGCTC CAAGTGCCAC TAACAATTCT TAGGACCTGA GCTATAAGCC
230881  AGGTGATTTC AGTTAATATG ATCAATTATT TCATTTAAAT GGCTCTAATG TGCAGAGGGA
230941  ACGGAGCCCA TCAGCATTCC CTGCAGGGAA CTGCAGTGGC TTTTATCAAC TTGAACAGCT
231001  AGCTTTCAAC TGTTTTGAAA TCACTTTCAG GGTGGTCATG TAGTTGCTTT TTTGAAATCA
231061  GAAGATGATT CTGCCTCTTT TAATATGTGA CTCCTCAGAT TCAGAAAGTG CTCGCTAGTC
231121  TTAAGAGTGA ATTACCCTCA GTGGTCCAGC GCTTATGAAC CCACATCTAA CCCTATCCCC
231181  TGGGGGAACT ATCAGAGAAA TTGGTGCCAT GGACATAAGA GGAAGGCACA GTGAAGCAGA
231241  GAGCCCCGCA TGATGAAAAT CAGTGGACAG CATCATTATT TACAACTTTG TAATCACCCA
231301  GGAGCATGAA AATCCAGGCC AATCTGGCAC CATGAGCTCT AATTTTTGTT GGAGTTCTTG
231361  GAACCGATTC TGATGAATGA CTGTTTAGCC ATTTTAGAGT GTGGCATACG TGGCTGCTGG
231421  CATACAGAGG TTGGATGTAA ACGGGCCTTT GCCCTCTCTT ATGAACATAG ACAGGAACTA
231481  AACTGTGTCA CATAGGTTCC AAATGGTGGC CTGAATACTA TTTACAACTA AGGTACAATG
231541  AAATTGAGTA AGTCTTTTCC TCTTTTGCAG ATACCATCAT TATTCATATA TTTCTTCAAA
231601  GTTAACTATT TGTATTTGGT AATTTTTAAT AGAAATGTAA TAATTGCTTC TCAAGTTTAG
231661  TCTTTAGTCT TAAGGTTGAT GCTCTCCATG TCCTTCCAAA AAAAGGTATG TTGCTTTTAT
231721  TATATCCTCG CCTTCAGATG GGATTATTCC ATTTTGTTCT TTGTTAATAT ATACTTTGAG
231781  CCACTTTTTT TGTGGCTCTG GGTGAGATGC TATAGGTACA ATGACAAGTG ATACGTGTGT
231841  TGTCCCTGTC ACAAAGTGG ATAGCCTAAG TGGTGACTTT TACCTCCACT CCAAATATAT
231901  GTATCACACA CCAGCCGTAT GCCAGGCACC ACTCTAGGTG CTAGGGATAC AGCAGTAAAC
231961  AGACAAATGC AACCCCTGCC CATGTGAAAG AGAATAAGAC AATAAATAAG TAAAGTGCAT
232021  GTTATATGGA GGTGGCAAAT GCTAAAAAGA AAAATTAAGC AGGCAAGAGG ACTCATTGAA
232081  AAGATGACAT TTGGGTAAAA GCCCATGTAT ATATGTTCTA TTGGTTTTAT TTCTCTGGAG
232141  AGCCCTGACT AATACACAAT GACTTTGAGA AGTTACTGGC TTTTGATTTA TCACACTATT
232201  CGGAGTGCTG AGAGCCTTCT TAGTGTGTAT TCAGTGTTTT AAGAGAGCTT GTGGATGAAT
232261  AATAAATAGG ACAAAATTTA TCCAAACTTA AGCCTTGCTT TAGGTAAAAG GGCTCCTCTT
232321  ACAAGGTAGA AGGTTATTAT TTGACATTTA AATCCAACTG AAGACTAATA AGACTAATTA
232381  ATTAAAAGTT TTTAAATCAC AACTGCGTGC AAAATAAATG GAACTGCCAT GCTCGCCAAG
232441  TGTGCATGAG TGGTGTGCAT GGGAGACAGC ACGAAGCTAA TCCCACTCAT CTTGCAGGTT
232501  GCTCCATTTT TCTCCTAAAA TCAGTAAGAC AGAAGCTGGT CAGATTATCA AGAGCCCTAG
232561  TTAAACACAG CAGTAGCATT TGGAAGGGGT TGCTCTCATT AGGCAGTGCC TGACCACAAC
232621  AAGAGATGAA CAAGCCCTGT ATCTGAAGCC ATCATGCCTA GTTATGGTCC CCGACTGTTC
232681  ATGATGCCTG GAAGGGAGGC CCCCTGCACC CTAGAAAGCT GGGTGGGTTC TACTGTCTGC
232741  TTTACTGCTA AAAACCCTCT TCTTTGGATC TGGACTTTAC CTCTATCTGA TTTTTTTTTC
232801  TAATATATGA TTTGGCACTG AGTCTGTCAC TGCTGCTAAC TCAGCAGTTC TAGGGTCATT
232861  GCCCCATTGC CTCACAGAAA GAATTTCATA GCTTCCAGCA TCCTCTCTCC TTCATTATAC
232921  TTTGATTTCA GCATTGCTAT TTTTTCTCTT GGGTGTTGCA GCTCTCTCTC TCCTTCCCAT
232981  GTCTTGTTGG TTTTCTGCTA ACTCCTGCTT TTTTTCTTTT TTTTTTTTTG AGACGGAGTC
233041  TCGTTCTGTC ACCCAGGCTG GAGTGCAGTG GCACAATCTC GGCTCACTGC AACCTCCGCC
233101  TCCCGGGTTC AAGCTATTCT CCTGCCTCAG CCTCCCAAGT AGCTGGGACT ACAGGCGCTC
233161  ACCACTATGC CCCACTAATT TTTGTATTTT TAGTATTGCT GTCATCAATC CACATGTCCA
233221  GAAGCACCTA GAAACTCTAA TTCTTTGTAG GTATCAAACC CTAGGACTCT TTCCTCTAAT
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233281   CACAATATAT AATCCCTGAT TCCCAAACAC GGTCTTTTCA TATACATTTT CCACTGTACA
233341   TACTTTCTGA CCTGGAAAGC TCTTACACAA ACACGCCCTC CCCTAGGAAG CCTTTATAAA
233401   TGTTCCCAGG AAGAATCAGT CACCCAACAG TGTCCTTGTC ACATCTTAGG TTCTACACCT
233461   TTATTTGTTC TATCTGAATG TAATCTCCCA GAGGGTGTTA TCATCTTTTT TTTTGAGATG
233521   GAGTCTTGCT TTGCTGCCCA GGCTGGAGTG CAGTGGCATG ATCTCGGCTC ACAGCAACCT
233581   CCACCTCCTG GGTTCAAGTG ATTCTCCTGC CTCAGCCTCC TGAGTAGCTG GGATTACAGA
233641   CGTGTGTCAC CACACCTGGC TAATTTTTGT ATTTTTAGTA GAGACAGGGT TTCACCGTGT
233701   TGGCAAGGCT TTCCTCGAAC TCCCAAACTC AGGTGATCCA CCCACCTCAG CCTCCCAAAG
233761   TGCTGGGATT ACAGGTGTGA GCCACCATGT CCAGCCCCAT CTTTTTCTTT TAGTTTAGTT
233821   CTTAACAAAT AGTCTGACAC AAAGTGGATA TAACAATATT TTGAATTATG AATAACTAAA
233881   TGAATATTTC CAGATTTCCT GGTGCTCTCA AAGTTTATG TTACAAAGA AAAACAAGTC
233941   TAAAATACCT GCCTCAAGTT TTTATCTGTA CTATGATTTC AAACCAAATA AAAAACAGGT
234001   GGGGTAAAAA CTGAAACAGG AAATACATAT AACTGAAAAA TTTTGGTATG TTAGTATGAT
234061   AATACTAGGT CATTTTTCCT GTTTCCCCAA CTTCATTTTC TATAGCAATA AAAAGAAACA
234121   AGTAAATGTA TGTTAATTTA ATTTAAAAGA AGTAGTCTAC CATCTCTTCT GTTAAAAAGA
234181   AAAAAGTATT TTAAAAAATT ATCTCTGGAA GGATACACAG GGAACATTGC TCTGGTTTCT
234241   TCCAAGAGAG AAATGAGGAA CTAGAGAGCA TGGCCAAGTG GGGTTTTGCT TTTGTTTTTG
234301   TTTGTCTATC TGTTAGCTTT TTATTATTTT CTTTTGTAGG TTTGAATTTC AAACCACATA
234361   AATCTGTTAC ATGCTCATAA TAATAAGTTT AAAATAAAAC TTTTGGCTGG GTGCAATGAC
234421   TTACACCTGT AATCCCAGCG CTTTGGGAAG CAGAGGTGGG AGGATACTTG AGGCCAGGAA
234481   TTTGAGATCA GCCTGGGCAA CATAGTGAGA CCCTGCCTCT GTAGAAATAA ACAAAAATTA
234541   GCTGGATATG GTGGTGCATG CTTGTACTCC TAGCTACTTG GGAGGTTGAG GCAGGAGGAT
234601   CCTTTGAGTC CAGGAGTTTG AGGCTGCAGT GAGCTATAAT CACCCACTGC ACTATAGCAT
234661   GGGCAATAAG GTGAGAACTT GTCTCAAAAA AAAAAGGGGG GGGGGAAACA AATAAATAAA
234721   TATAAACAAA ACTTTTGTTT CAAAATATGT AATATTTAGC ACTAAAGAAT TCTGAATTGT
234781   AGAGCTAAAA AGTACTTAAA AGTTAATAAC TATTGTCTCC TTTAAAAGAA TTGTTATCAA
234841   AGTATAATTT TTATCCAGAA AATCATCCAT ATCAGCAAGC TAAACTTTCT CAAAATGACA
234901   TATCCATGTA ATTAGCTCCC AGGTAATTAG CAGGCAGCCT CTACTCAGGT TGAGTATTCC
234961   TAATCTAAAA ATTGGAAATT CAAAATGCTC CAAAATCTGC AACTTTTTGA ATGCTAACAT
235021   GATTCTCAAA GGAGTGCTCA TGGAGTATTT CAGATTTTGG ATTTTTGGAT TTGAGATACT
235081   CAGTATAATG CAAACATTCC AAATCTGAAA AAATCTGAAA TACTTCTGGT TCTAAGCATA
235141   AGGGATACTC AACGTGTGTT AGCTAATTAG ACCCTTCATG GTCTCTTCTA GACCTCAGCT
235201   TCTTCAAGGT AACCTCTATC CTCACTTCTA ATAGCATGAA CTTTTCTGTT TTAGAATAAT
235261   TTGGATTTTC AGGAAAGTTG CAAAGATAGT ACAAAGACAG TACAGGAGAG TTCCCATATA
235321   TCTTTCACCT AGCTTTCCCC CATTGTTAGG ATTTTACATT ATTATGATAC ATTTGTCAAA
235381   TATAAGCAAC TCACATTGAT ACATGAAACT CTATTAACCA AACCCTAGAC TTTATGTGGA
235441   TTTCACCACT GTTCCACTA ATGTTTTCTT TCTGTTCCAA GGTCCAATCT GGAATACCAC
235501   ACTGCATTTT CTTGTCATAT CTCCCTAGTC TTTTTTTGTC TGTGACAATG TCTCAGTCTT
235561   TTCTTGCTTT TCATGACCTT AACAGTCCTG AAGATCATTT GCTTTTTTTT CATAATTACA
235621   CCGGAGTTAT AGATTTTTTG AAATAATACC ACAAGGGCAA AGGGCCCTTC TTGTCACATC
235681   ATTTTAGGGA GAACATGATA TCCACATGAC ATCACTGATA TTAACCTTCA TCATGTGGTT
235741   TAGGTAATGT TTCAGGTTTC TCTACTGCAA AGTGATTTTT TTCCCTTAAT TTAGCCCACC
235801   TGAACTTATC AATTTTGTTT TCTTCCATGA CTAATACTTT TGTTATTATA GCTAAAACTT
235861   CATTGGGGCC AAATCTTAGA TCATGTAAAT TTTCTTCTAT ATTTTATTCT AAAAGCTTGT
235921   AATGTTTGAT ACATTCTAAA AGATGTAATG TTTGATACAT TACATCTAGT CCTTTGATTT
235981   ATTTTTAGTT ACTTTTGTAT AAGGTGTGAG AGATGTCTCC AGTTTCACTT TATTAACACA
236041   TTGTGGTGTT CCAGTACTAT TTGTTGCTAA GACTATCTTT TTTCCATTGA TTACCTTTGC
236101   CTTAGTTGGC AATATTTTTG TTGGTTTATT TCTAGACTGT TTATCTCATT CCACTGATTT
236161   GTGTCTATCT TTTTGACAAA ACTGTTGATT ACAGTAAGCT TTGAAATAGT TCATTTTTTG
236221   TGTCAACTTG ACTGAGTCAG GGGATAACCA GCTATCTGGT TAAACATTAT TTCTGGCTGT
236281   GTTTGTGAGC GTGTTTCTGG ATGAGATTAG CCTTTGAATA GGTGATCCTA GTAAAGTAAA
236341   CTGTCTTTCC CAGTGTGGAT GGCATTATGC CACCTGATAT TCAGGGTCTG AATAGAAGAA
236401   AAGGCAGAGG AAGGGGGAAT TTGGGCCTTT TTTTCTGCCT CACTGCTTGA GCTGGGACAT
236461   CTCATCTGGT CTCCTGCTCT TGAACTGGGA TTTACATCAT CAGTTCCTCT GGTTCTCAGG
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236521   CCTTCAGATT CAGACTGAAT CATACCACCA GCTTTCCTGG GTCTCCAGCT TGCAGATTAC
236581   AGATCATGGG ACTCCTCATC TTCCATAAAT GCATGAGCCA ATTCAGTCTA TGTCCTTGAA
236641   AACTGCCCCA CTGCAGATTA AGGCTTTTTT CCACTAGGTG AAATAAAGAA GCTTGTTAGA
236701   CAGATTTCCC TTCATCCAGT GCCCTCTCCT CTTTAAGTTA CAACACATTG GCTACACCTA
236761   AGTGCAGGGG TGGGGATGAG GGTATAGTCC TCTTGTTTGC TGAGAAGAGA ACTGTATTGG
236821   GAAAGCTCTA GAAGTGTTTG ATACATACAT AAACAAGGCA TGGTTTTTGC ACTTAATTTC
236881   ACATTACATT TTTCCCAGAA AAAAAGGAAT GTATAGGCAT CACGTAACTG TACTAGCTGG
236941   AGTCATTCTT CCTGATTATC AAAGGTAAAC AGTTATTAAT CCTATACCAA GATGTCAAGG
237001   AGAAGTACTT TTGGAACACA AGGAATTCTC TGGGAGTCCT TACTACTCTC AAGCCCAGTG
237061   AAAAAGTTAA TGAAAAACTA TAGTACCTTC CTATAAGCTG GATGACTAAT TACCAGGCTC
237121   ATTTAGGAAT TTGCCTTACC AAGTAAAACA TAAGGGCAGC TGAGGTGCTG ACTGAAGACA
237181   AATGGAGCAT AGAATAAGAG TAGTAAAGAA TGCCAAAAAT GCTGTCATGT ATCCATTGAC
237241   AAAAGGAGCT ATAAAGCCTT TAGGTATTTT CACACTTGCT CTGTTACGTA AATGTATGTG
237301   TGTGTGTGTG TGTGTGTGTG TGTGTG
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A. CLASSIFICATION OF SUBJECT MATTER

IPC(6) : C07H 21/04; C12Q 1/68; C12N 15/63, 15/85; C12P 21/02
US CL : 536/23.5; 435/6, 70.1, 325, 320.1
According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 536/23.5; 435/6, 70.1, 325, 320.1

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

APS, DIALOG'S BIOTECH cluster.
hemochromatosis, BTF1, BTF2, BTF3, BTF4, NTP-3, NTP-4, RoRet, butyrophilin, type 1 sodium transport

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A, P	RUDDY, D.A. et al. A 1.1-Mb transcript map of the hereditary hemochromatosis locus. Genome Research. May 1997, Vol. 7, No. 5, pages 441-456, see entire document.	1-20, 22-77
X	FISCHER, L. et al. Cloning of the 62-kilodalton component of basic transcription factor BTF2. Science. 04 September 1992, Vol. 257, pages 1392-1395, see entire document.	28-33, 71
X	MARGOTTIN, F. et al. Participation of the TATA factor in transcription of the yeast U6 gene by RNA polymerase C. Science. 25 January 1991, Vol. 251, pages 424-426, see entire document.	22-27, 70

X	Further documents are listed in the continuation of Box C.		See patent family annex.

	Special categories of cited documents:	"T"	later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
"A"	document defining the general state of the art which is not considered to be of particular relevance		
"E"	earlier document published on or after the international filing date	"X"	document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
"L"	document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	"Y"	document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
"O"	document referring to an oral disclosure, use, exhibition or other means		
"P"	document published prior to the international filing date but later than the priority date claimed	"&"	document member of the same patent family

Date of the actual completion of the international search	Date of mailing of the international search report
20 JANUARY 1998	1 2 FEB 1998

| Name and mailing address of the ISA/US
Commissioner of Patents and Trademarks
Box PCT
Washington, D.C. 20231	Authorized officer F. PIERRE VANDERVEGT
Facsimile No. (703) 305-3230	Telephone No. (703) 308-0196

C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	ZHENG, X.M. et al. Sequencing and expression of complementary DNA for the general transcription factor BTF3. Nature. 05 April 1990, Vol. 344, pages 556-559, see entire document.	34-39, 72
X	PANTEGHINI, M. Electrophoretic fractionation of 5'-nucleotidase. Clinical Chemistry. February 1994, Vol. 40, No. 2, pages 190-196, see entire document.	52-57, 75
X ---- A	BURT, M. J. et al. A 4.5-megabase YAC Contig and physical map over the hemochromatosis gene region. Genomics. 15 April 1996, Vol. 33, No. 2, pages 153-158, see entire document.	1-6 ---- 7-20, 22-77
A	VERNET, C. et al. Evolutionary study of multigenic families mapping close to the human MHC Class I region. J. Mol. Evol. November 1993, Vol. 37, No. 6, pages 600-612, see abstract in particular.	1-20, 22-77

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US97/17658

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This international report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:

2. ☐ Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:

3. ☐ Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

Please See Extra Sheet.

1. ☒ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.

2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.

3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:

4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest   ☐ The additional search fees were accompanied by the applicant's protest.

               ☒ No protest accompanied the payment of additional search fees.

Form PCT/ISA/210 (continuation of first sheet(1))(July 1992)★

BOX II. OBSERVATIONS WHERE UNITY OF INVENTION WAS LACKING
This ISA found multiple inventions as follows:

This application contains the following inventions or groups of inventions which are not so linked as to form a single inventive concept under PCT Rule 13.1. In order for all inventions to be searched, the appropriate additional search fees must be paid.

Group I, claim(s)1-20, drawn to polynucleotide sequences containing at least one polymorphic site, polypeptides encoded thereby, antibodies to said polypeptides and a method to determine the presence of the HFE gene mutation.

Group II, claim 21, drawn to the lymphoblastoid line atcc crl-12371.

Group III, claim(s) 22-27 and 70, drawn to BTF1 nucleic acids, gene products, vectors and antibodies.

Group IV, claim(s)28-33 and 71, drawn to BTF2 nucleic acids, gene products, vectors and antibodies.

Group V, claim(s) 34-39 and 72, drawn to BTF3 nucleic acids, gene products, vectors and antibodies.

Group VI, claim(s) 40-45 and 73, drawn to BTF4 nucleic acids, gene products, vectors and antibodies.

Group VII, claim(s) 46-51 and 74, drawn to BTF5 nucleic acids, gene products, vectors and antibodies.

Group VIII, claim(s) 52-57 and 75, drawn to NPT3 nucleic acids, gene products, vectors and antibodies.

Group IX, claim(s) 58-63 and 76, drawn to NPT4 nucleic acids, gene products, vectors and antibodies.

Group X, claim(s) 64-69 and 77, drawn to RoRet nucleic acids, gene products, vectors and antibodies.

The inventions listed as Groups I-X do not relate to a single inventive concept under PCT Rule 13.1 because, under PCT Rule 13.2, they lack the same or corresponding special technical features for the following reasons: Groups I and III-X are drawn to physically different genes and their gene products and each therefore constitutes a separate invention. The lymphoblastoid cell line of Group II is not dependent upon the vectors of any of the Groups I and III-X and therefore constitutes a separate invention. Accordingly, the claims are not so linked by a special technical feature within the meaning of PCT Rule 13.2 so as to form a single inventive concept.